
CPN Workshop `04

Application of Coloured Petri Nets in System Biology

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Germany**

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CPN Workshop`04 – Application of Coloured Petri Nets in System Biology

**main aim: model development of metabolic networks,
 which allows validation techniques**

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validation techniques:

T-invariant analysis

➤ introduced by Heiner et al. [1]

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T-invariant analysis

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P-invariant analysis

- **P-Invariant corresponds to mass conservation law**
 - ⇒ covering P-invariant necessary ⇒ bounded net
- **inconsistencies between reaction & compound formulas detectable**

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validation techniques:

T-invariant analysis

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P-invariant analysis

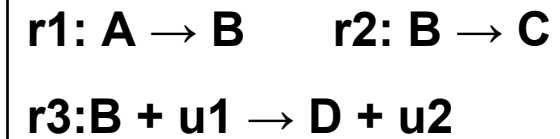
- P-Invariant corresponds to mass conservation law
 - ⇒ covering P-invariant necessary ⇒ bounded net
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model checking

- more detailed prove of properties / conditions
 - ⇒ requires finite state space

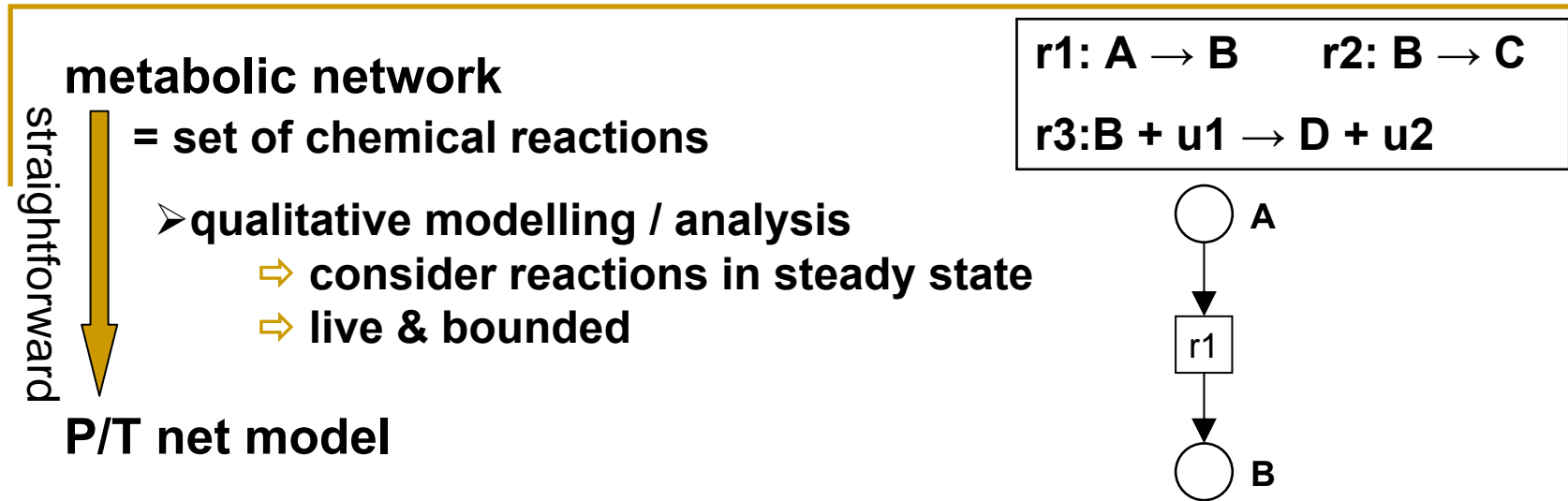
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metabolic network
= set of chemical reactions

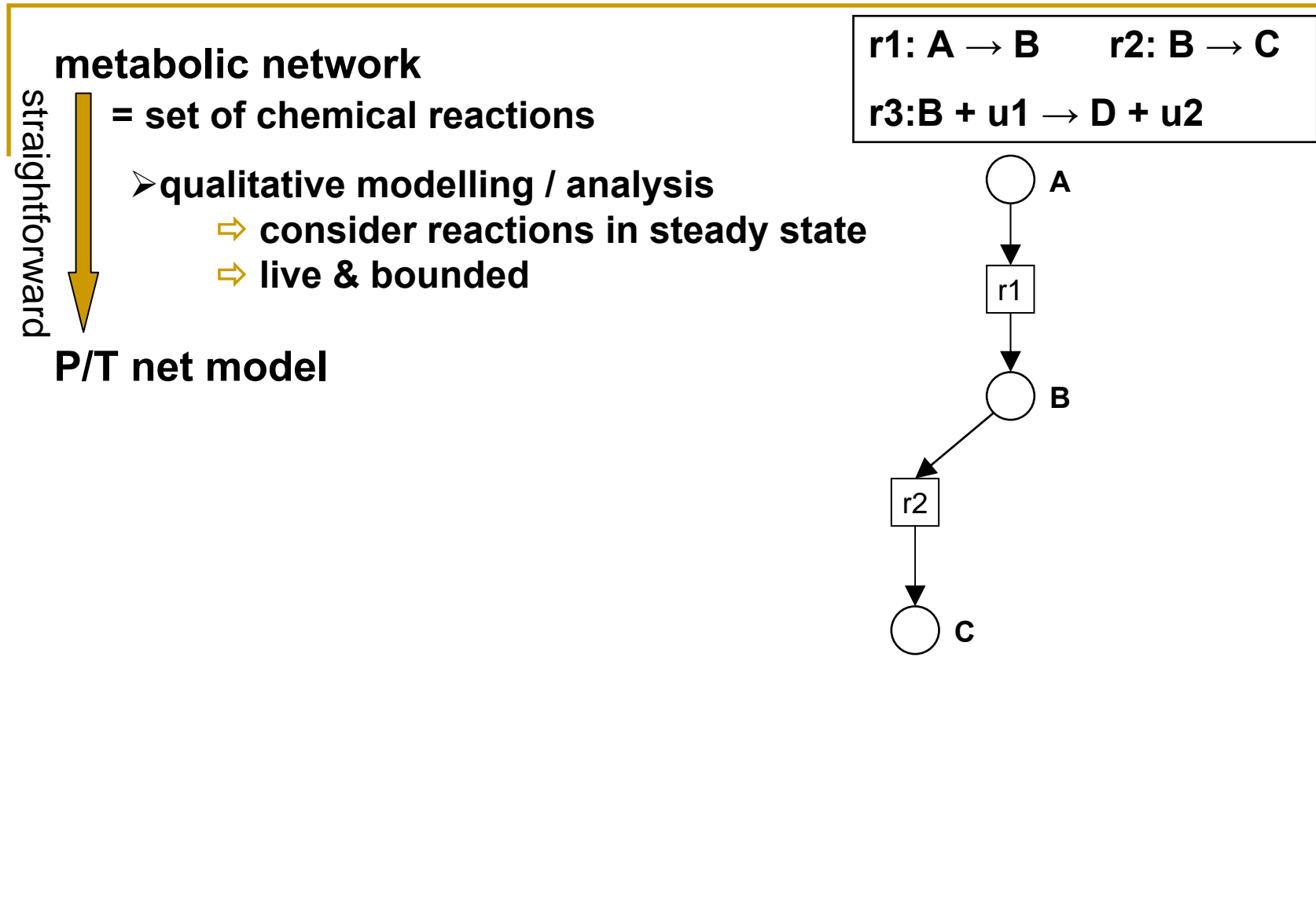


- **qualitative modelling / analysis**
 - ⇒ consider reactions in steady state
 - ⇒ live & bounded

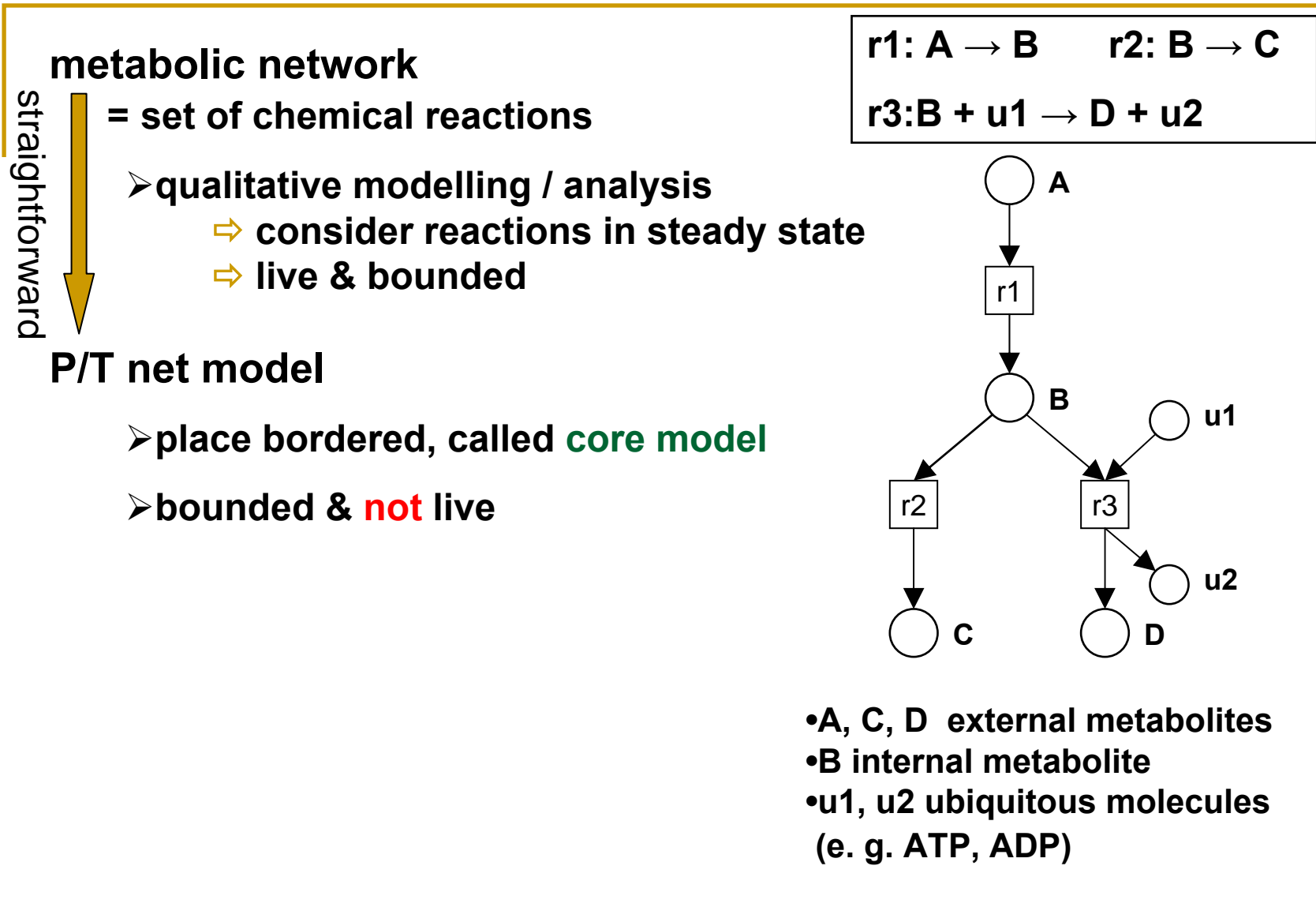
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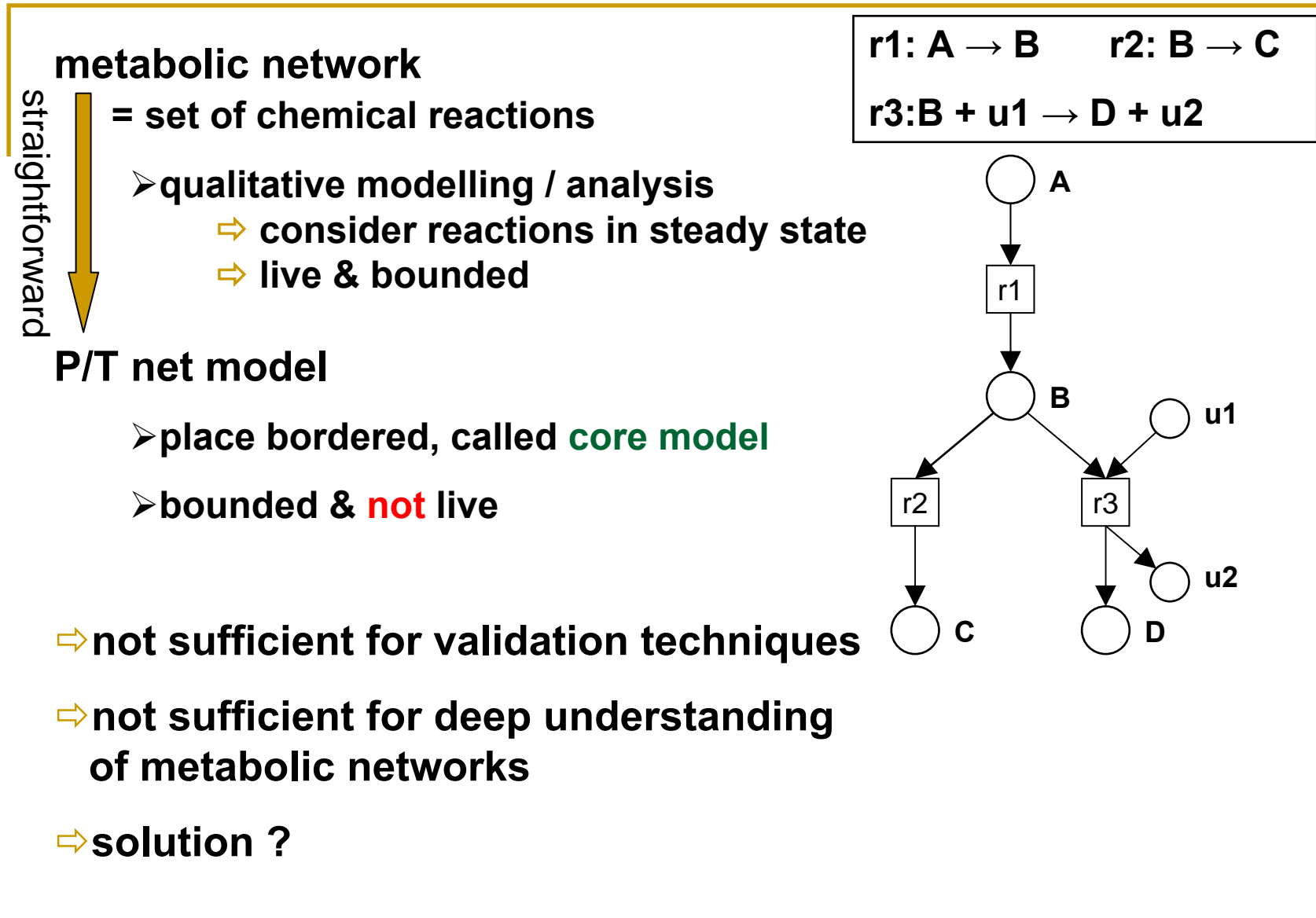
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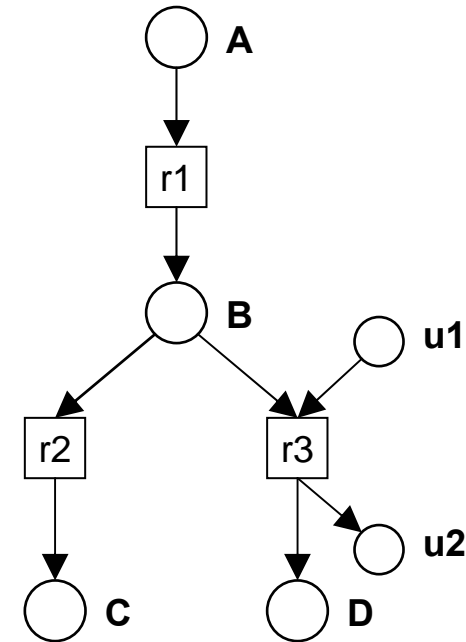
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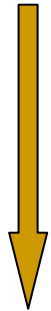
P/T net model (bounded & not live)

- requires environment behaviour
- easiest biochemical behaviour
 - ⇒ supply & removal



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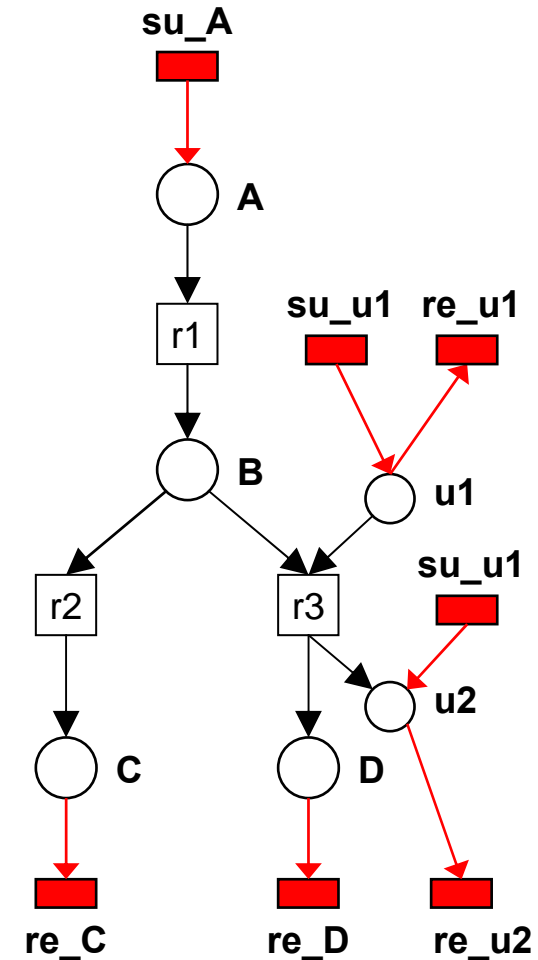
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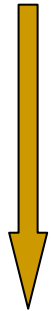
P/T net model with environment type I

- transition bordered, open system,
called **system model (I)**



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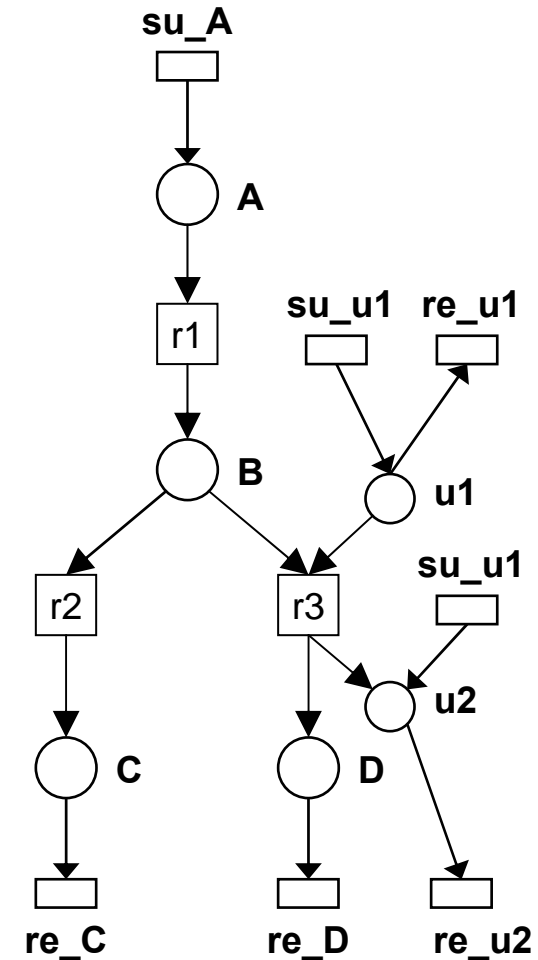
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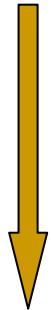
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- **un**bounded & live & covered by T-invariants (CTI)



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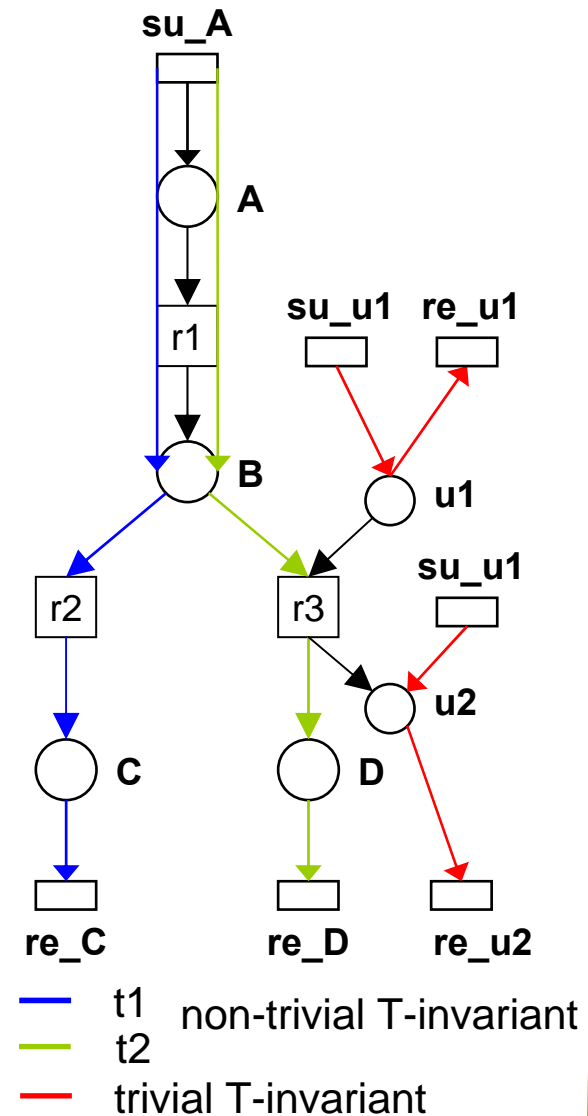
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P/T net model with environment type I

- transition bordered, open system, called **system model (I)**
- **un**bounded & live & covered by T-invariants (CTI)
- minimal T-invariant = elementary mode (biochemistry)
⇒ deep understanding (pathways)



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P/T net model (bounded & not live)



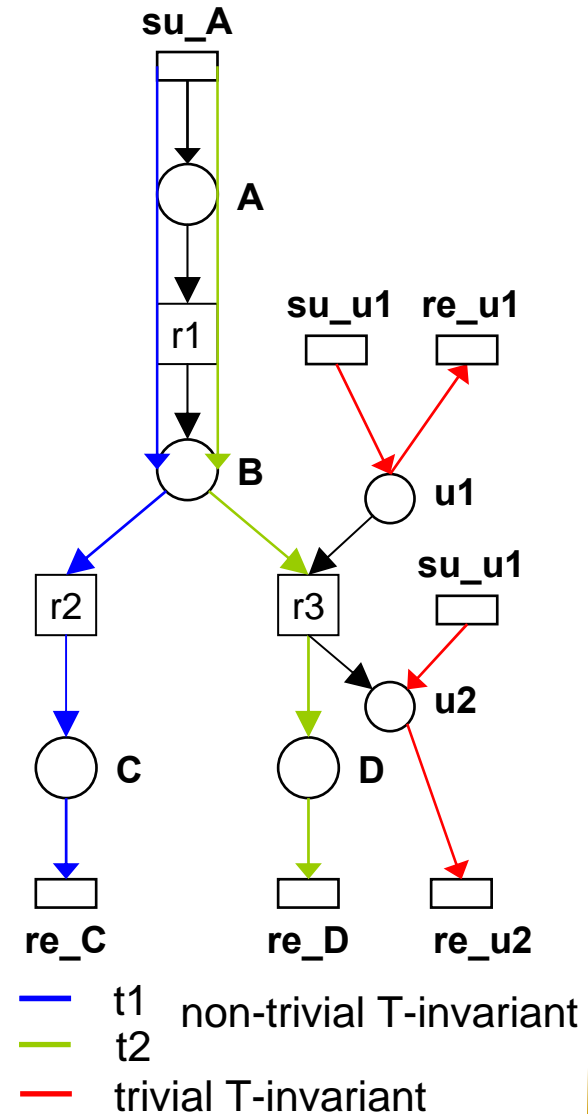
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⇒ **P-invariant analysis ?**

⇒ **model checking ?**



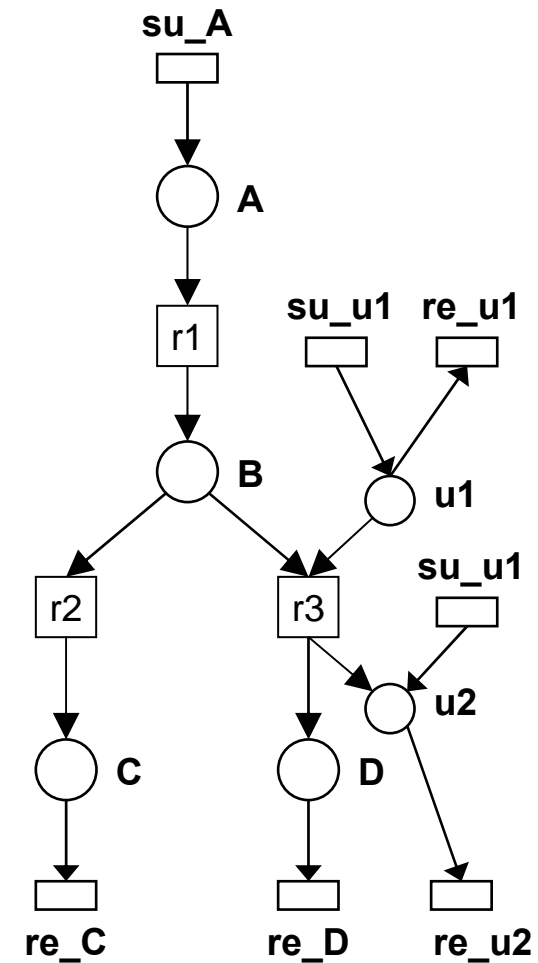
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P/T net model with environment type I



CPN model with environment type II

- closed system,
called **system model (II)**



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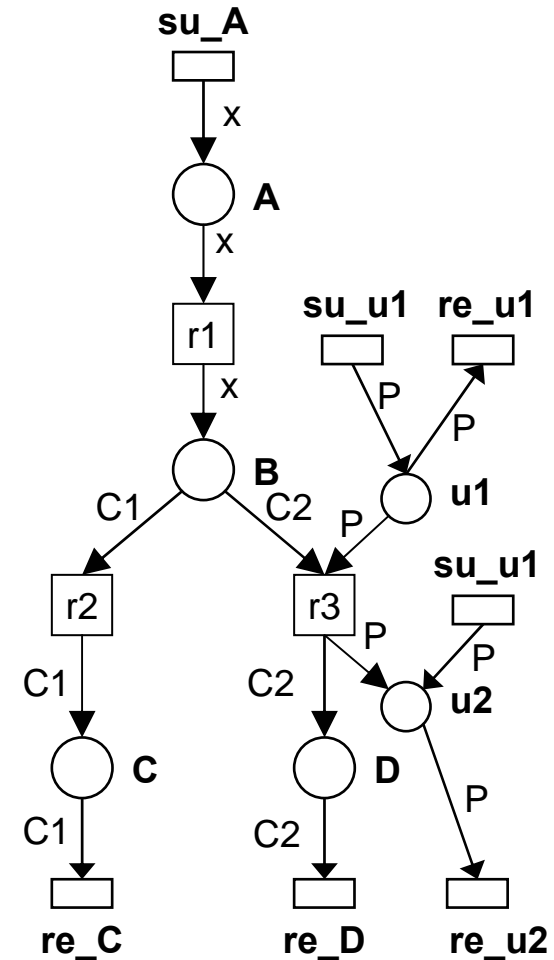
P/T net model with environment type I



- colour the model to avoid dynamic conflicts (dead markings)
- ⇒ two variants

CPN model with environment type II

- closed system, called **system model (II)**



color CS = C1 | C2 | P;
 color X = subset CS with [C1, C2];
 var x:X;

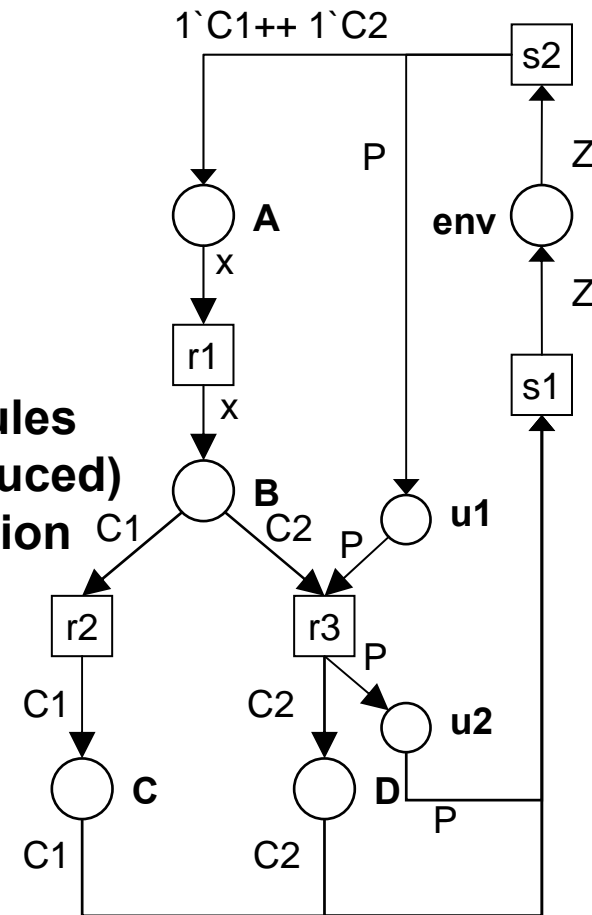
P/T net model with environment type I

construction

- colour the model to avoid dynamic conflicts (dead markings)
 - ⇒ two variants
- transform environment behaviour
 - ⇒ limit amount of suppliable molecules
 - ⇒ supply (remove) consumed (produced) molecules by a start (stop) transition

CPN model with environment type II

- closed system, called **system model (II)**



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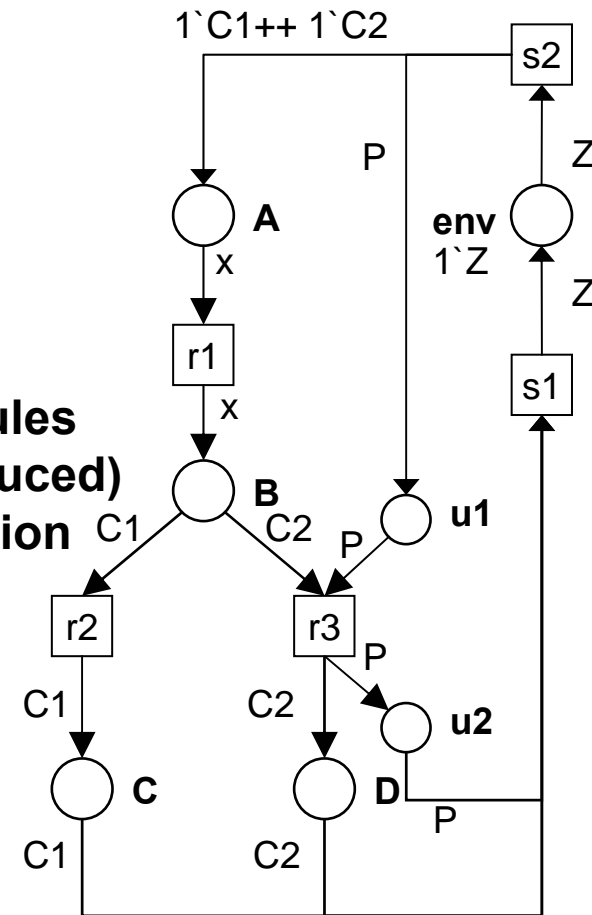
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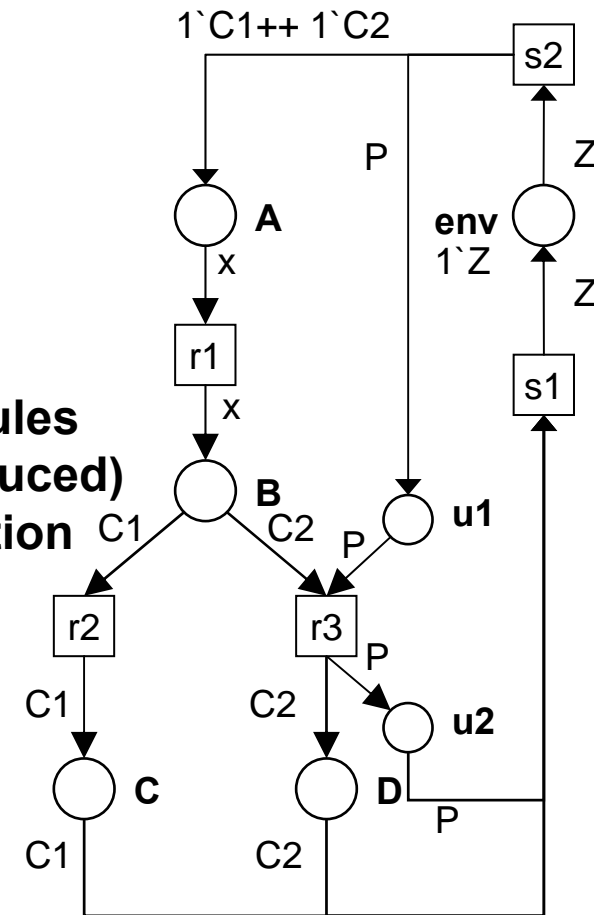
P/T net model with environment type I



- colour the model to avoid dynamic conflicts (dead markings)
 - ⇒ two variants
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CPN model with environment type II

- closed system, called **system model (II)**
- bounded & live & CTI & covered by P-invariants (CPI)
- only small subset of CPN used



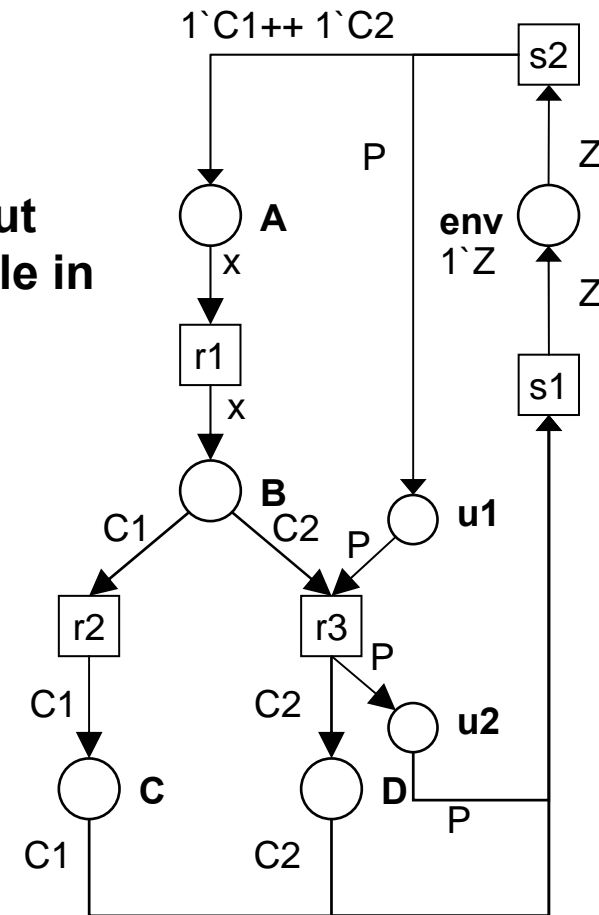
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P/T net model with environment type I

↑ behavioural equivalence

- Each realizable T-invariant of the unbounded system model (I) without border transitions must be realizable in the bounded system model (II).

CPN model with environment type II



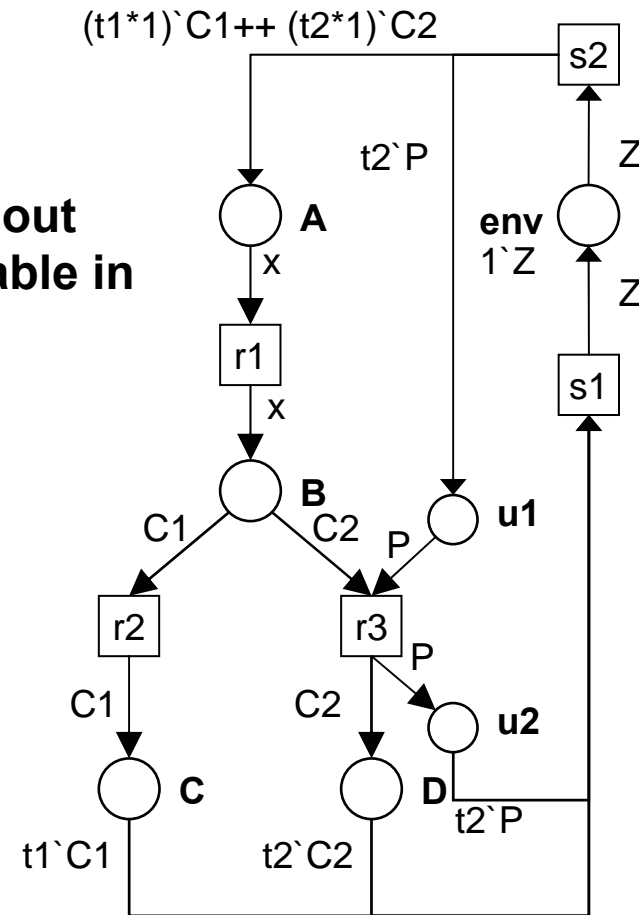
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P/T net model with environment type I

↑ behavioural equivalence

- Each realizable T-invariant of the unbounded system model (I) without border transitions must be realizable in the bounded system model (II).
- normally fusion of T-invariants
- selection parameter (t_1 and t_2)

CPN model with environment type II



color CS = C1 | C2 | P | Z;
 color X = subset CS with [C1, C2];
 var x:X; val t1 = 1; val t2 = 1;

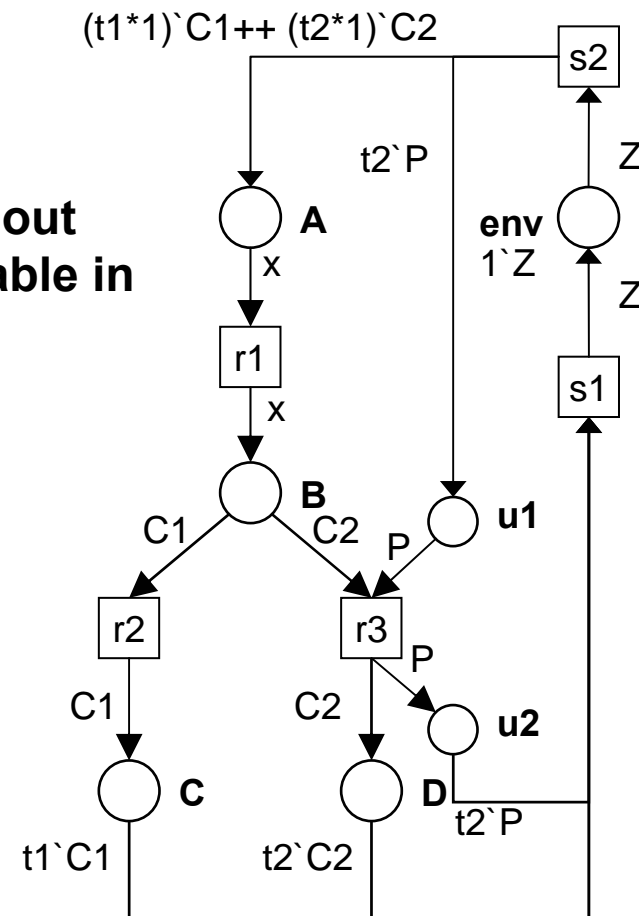
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CPN model with environment type II

- CTI & CPI
 - ⇒ prove using effect / defect calculation²
- bounded & live
 - ⇒ prove using Occ/ Scc graph calculation



color CS = C1 | C2 | P | Z;
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CPN model with environment type II

➤ CTI & CPI

⇒ prove using effect / defect calculation

⇒ multiplying $P \times T$ -incidence matrix C with transformation vectors

effect

$C \cdot x = z$; x .. substitution vector (T-vector),
 z .. **effect of x** = P-vector, which denotes marking differences for each place

$z = 0$ ⇒ x is T-invariant

substitution ⇒ **binding element**

e. g. $r1: x \leftarrow (1 \text{ } C1), (1 \text{ } C2)$

CPN model with environment type II

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effect - example

	r1	r2	r3	s1	s2
A	-x				C1+ C2
B	x	-C1	-C2		
C		C1		-C1	
D			C2	-C2	
u1			-P		P
u2			P	-P	
env				Z	-Z

incidence matrix C

*

r1: $x \leftarrow (1 \text{`C1}), (1 \text{`C2})$
r2: $_ \leftarrow (1 \text{`_})$
r3: $_ \leftarrow (1 \text{`_})$
s1: $_ \leftarrow (1 \text{`_})$
s2: $_ \leftarrow (1 \text{`_})$

substitution vector

= 0

⇒ T-invariant

⇒ CTI

* replaces variables by expressions in token colours

CPN model with environment type II

➤ CTI & CPI

⇒ prove using effect / defect calculation

⇒ multiplying $P \times T$ -incidence matrix C with transformation vectors

defect

$C^T \cdot y = w$; y .. distribution vector (P-vector),
 w .. **defect of y** = T-vector, which denotes marking differences caused by each transition

$w = 0$ ⇒ y is P-invariant

sensible biochemical distributions

e. g. Glucose $C_6H_{12}O_6$ ⇒ Gluc: $\rightarrow 6`C + 12`H + 6`O$

CPN model with environment type II

➤ CTI & CPI

⇒ prove using effect / defect calculation

⇒ multiplying $P \times T$ -incidence matrix C with transformation vectors

defect - example

	A	B	C	D	u1	u2	env
r1	-x	x					
r2		-C1	C1				
r3		-C2		C2	-P	P	
s1			-C1	-C2		-P	Z
s2	C1+ C2				P		-Z

transposed incidence matrix C

$$\begin{array}{l}
 \text{A: } C1, C2, x \rightarrow (1^{\setminus}C1), (1^{\setminus}C2), (1^{\setminus}C1 + 1^{\setminus}C2) \\
 \text{B: } C1, C2, x \rightarrow (1^{\setminus}C1), (1^{\setminus}C2), (1^{\setminus}C1 + 1^{\setminus}C2) \\
 \text{C: } C1 \rightarrow 1^{\setminus}C1 \\
 \text{D: } C2 \rightarrow 1^{\setminus}C2 \\
 \text{u1: } P \rightarrow 1^{\setminus}P \\
 \text{u2: } P \rightarrow 1^{\setminus}P \\
 \text{env: } Z \rightarrow 1^{\setminus}C1 + 1^{\setminus}C2 + 1^{\setminus}P
 \end{array}
 = 0$$

* transforms elements of colour set D into linear combination of elements of set E

distribution vector (imagined)

⇒ **P-invariant**

⇒ **CPI**

CPN model with environment type II

- bounded & live
- ⇒ prove using Occ/ Scc graph calculation
- artificial example CPN model (not shown)

	places	transitions	token colours	T-invariants	Occ graph	
					nodes	arcs
variant I	10	10	5	4	92.177	513.047
variant II			8		274.187	1.707.422

- case study: extended glycolysis
- ⇒ 33 places, 21 transitions and 40 T-invariants
- ⇒ Occ graph calculation not completed (limited resources)

Summary

- method to construct behavioural equivalent system model (II) from system model (I) using coloured Petri nets
⇒ validation possible
- verification of P- and T- invariants for subset of coloured Petri nets
- conditions to prove with model checking? ⇒ biochemists

References

- ¹ HEINER, M. ; KOCH, I.: Petri Net Based Model Validation in Systems Biology. In: *LNCS* Bd. 3099, Springer, 2004, S. 216–237
- ² VOSS, K. ; HEINER, M. ; KOCH, I.: Steady state analysis of metabolic pathways using Petri nets. *In Silico Biol.* 3 (2003), Nr. 3, S. 367–387