

PETRI NETS 4 BACTERIAL BIOENGINEERING

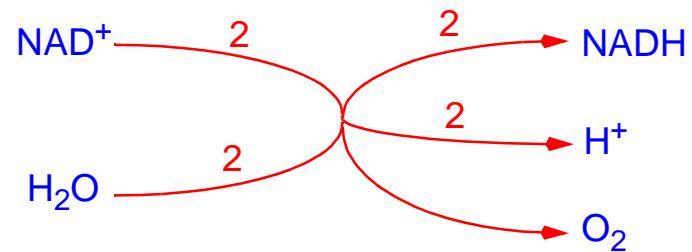
Monika Heiner

*on sabbatical leave from Brandenburg Technical University
Computer Science Institute*

THE PETRI NET FRAMEWORK

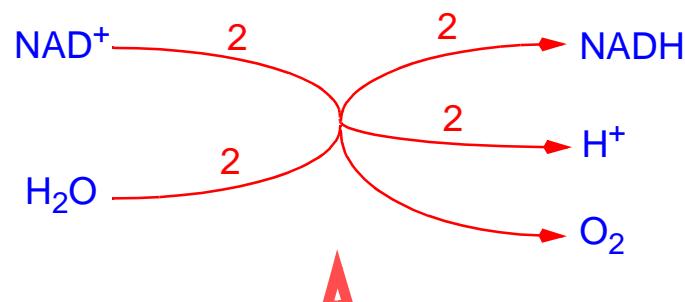
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ARE NETWORKS OF BIOCHEMICAL REACTIONS

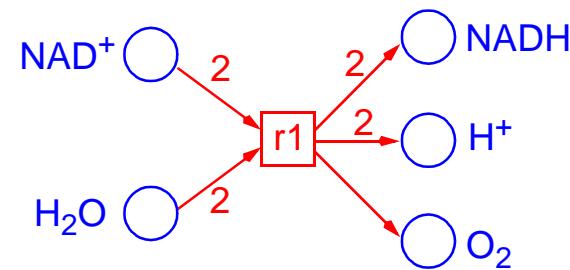


ARE NETWORKS OF BIOCHEMICAL REACTIONS

NATURALLY EXPRESSIBLE AS PETRI NETS

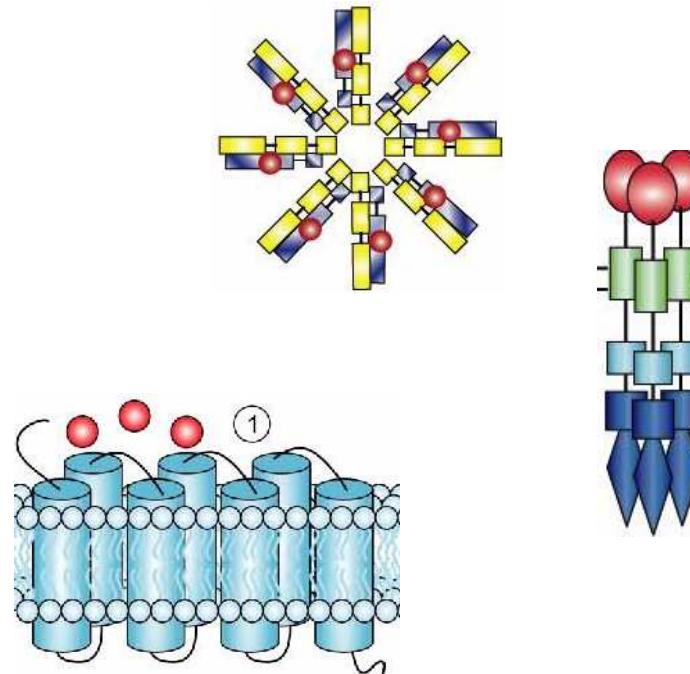


hyper-arcs



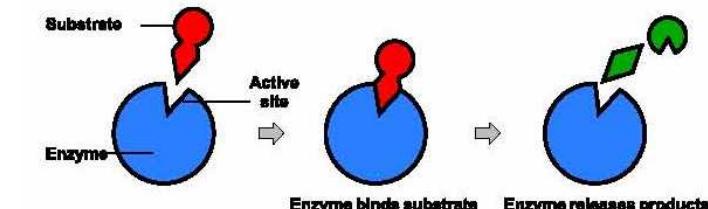
□ places → model variables

- > (bio-) chemical compounds
- > proteins
- > protein conformations
- > complexes
- > genes, . . . , etc.
- ... in different locations*

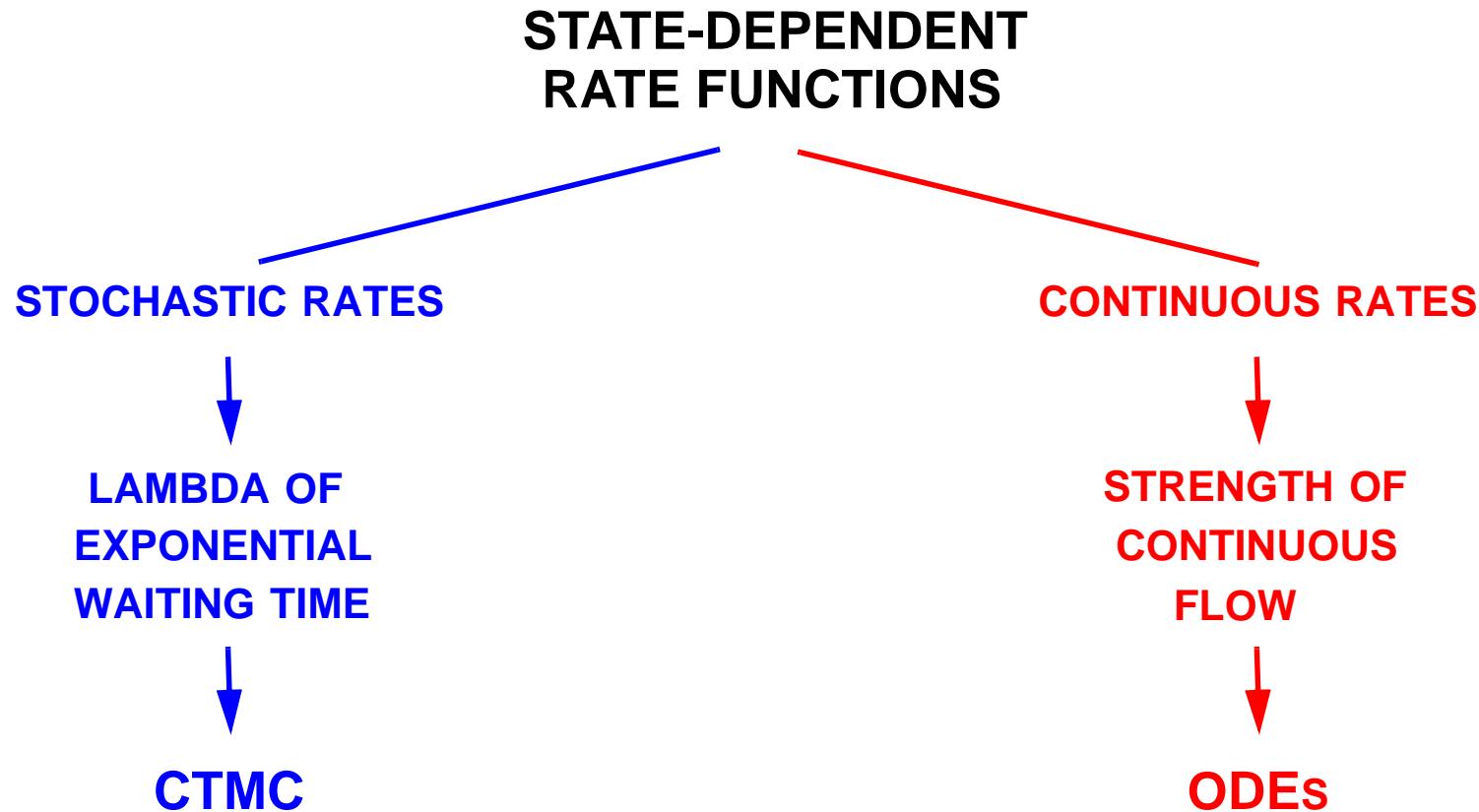


□ transitions → atomic events

- > (stoichiometric) chemical reaction
- > complexation / decomplexation
- > phosphorylation / dephosphorylation
- > conformational change
- > transport step, . . . , etc.
- ... in different locations*



STATE-DEPENDENT RATE FUNCTIONS

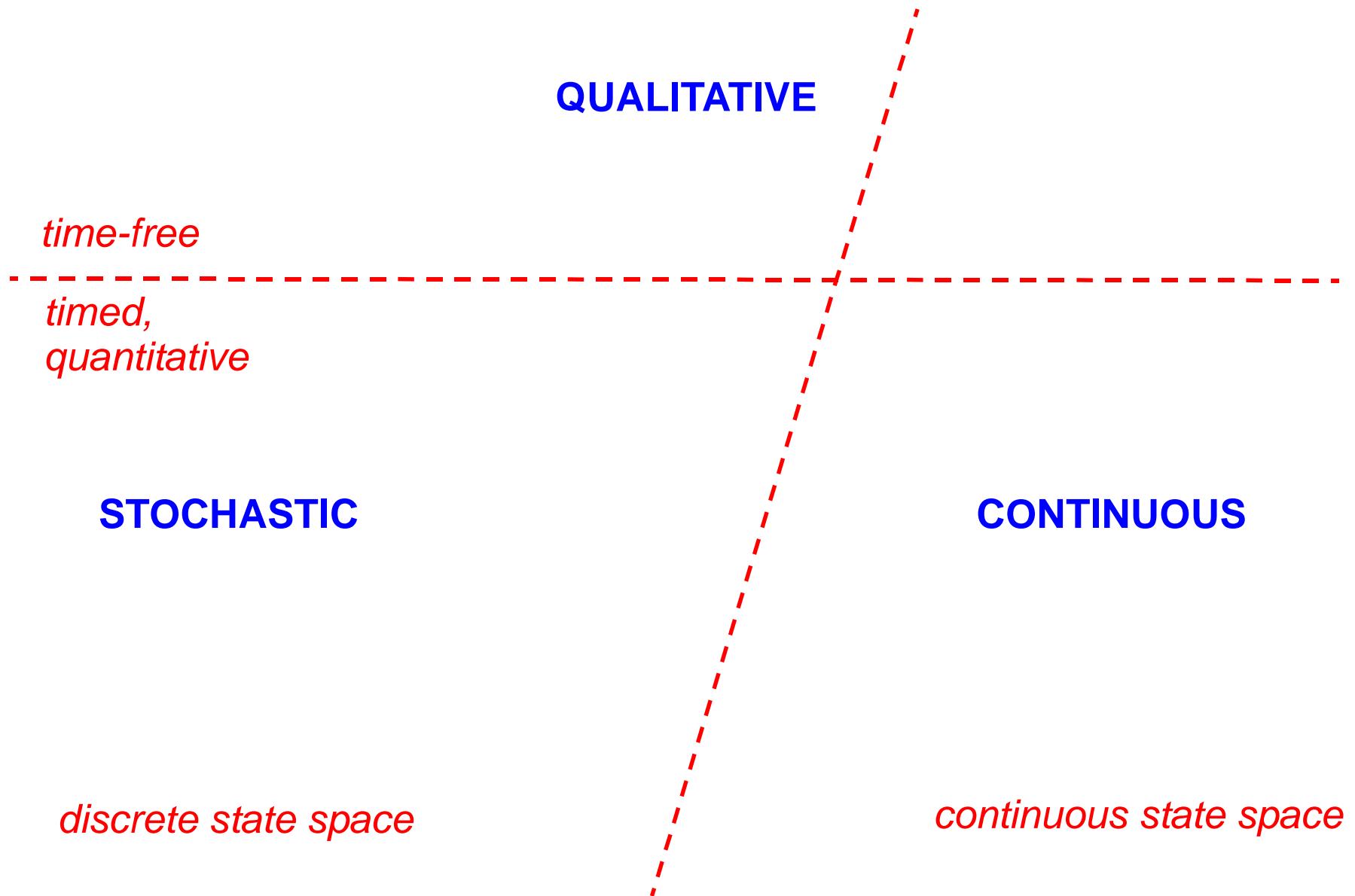


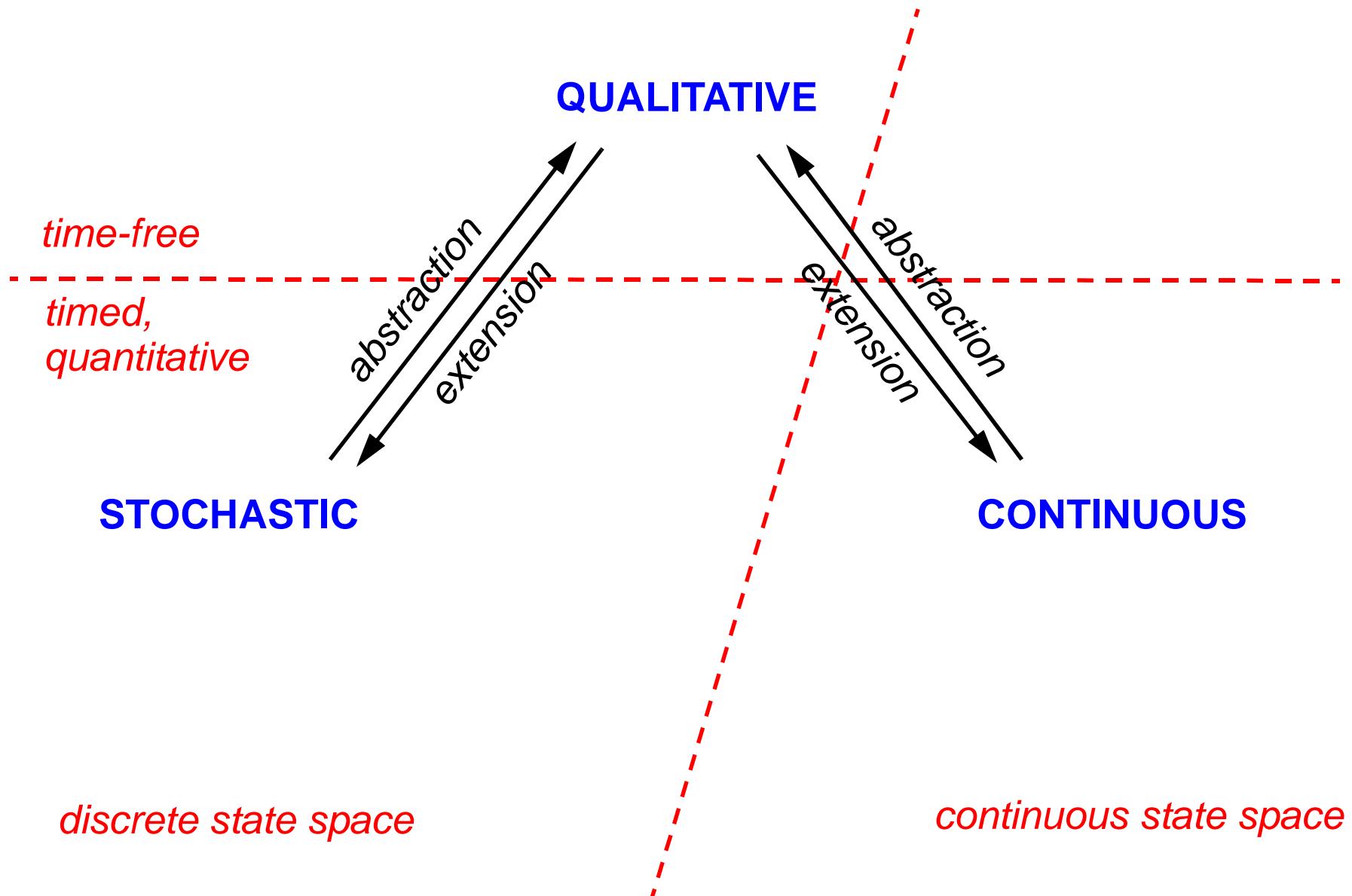
-> supported by, e.g., COPASI, Dizzy, ..., Snoopy

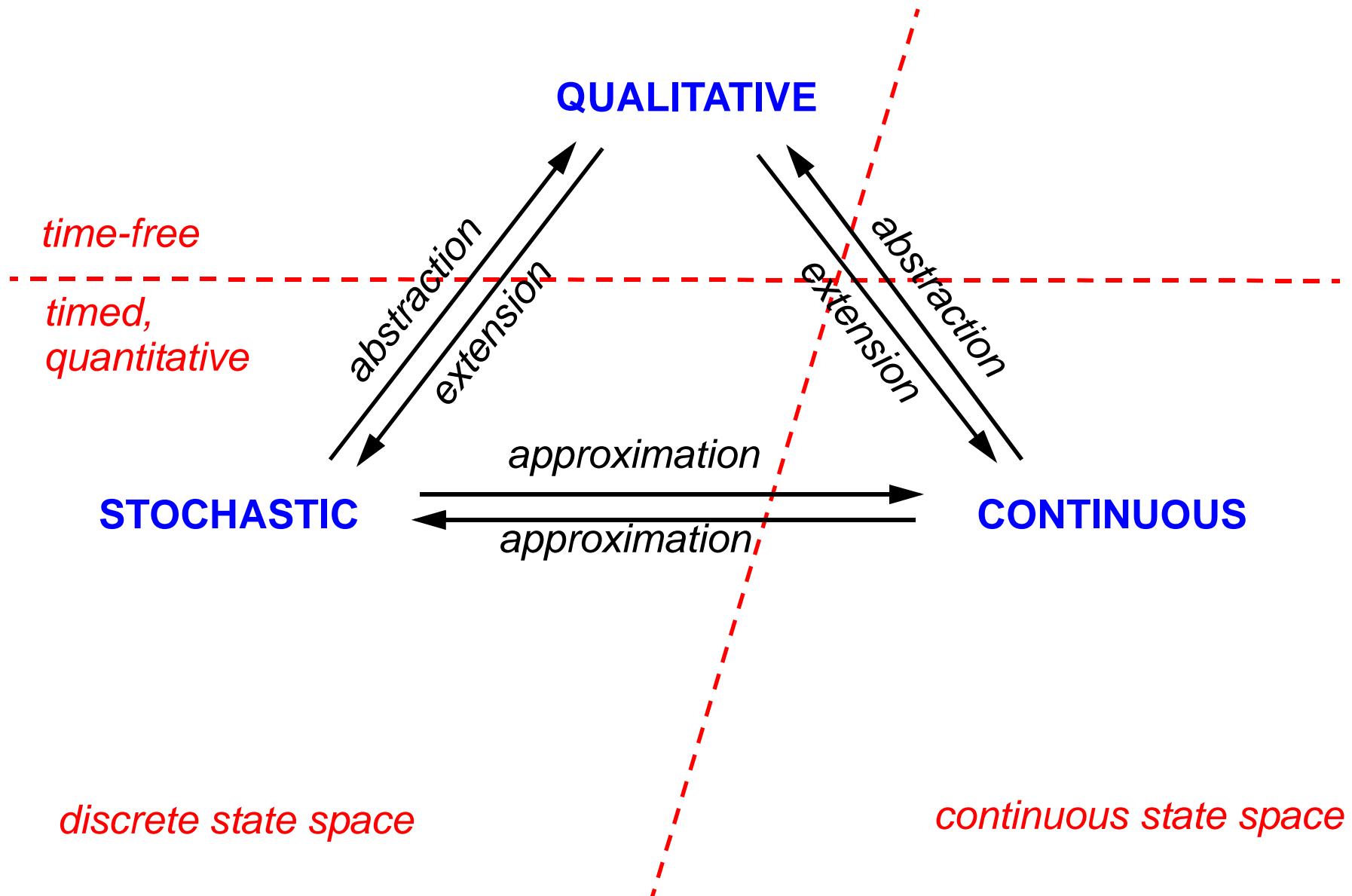
QUALITATIVE

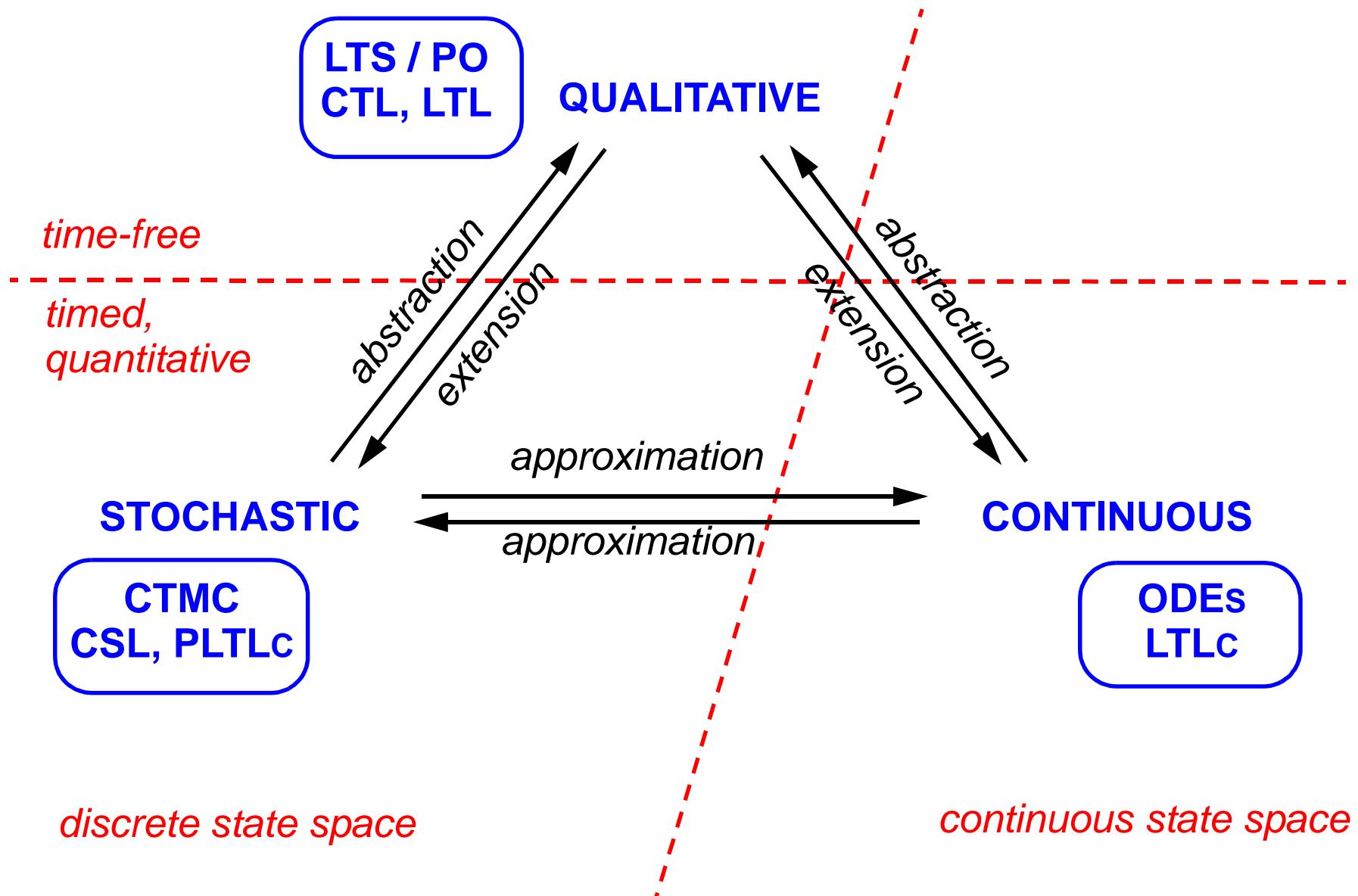
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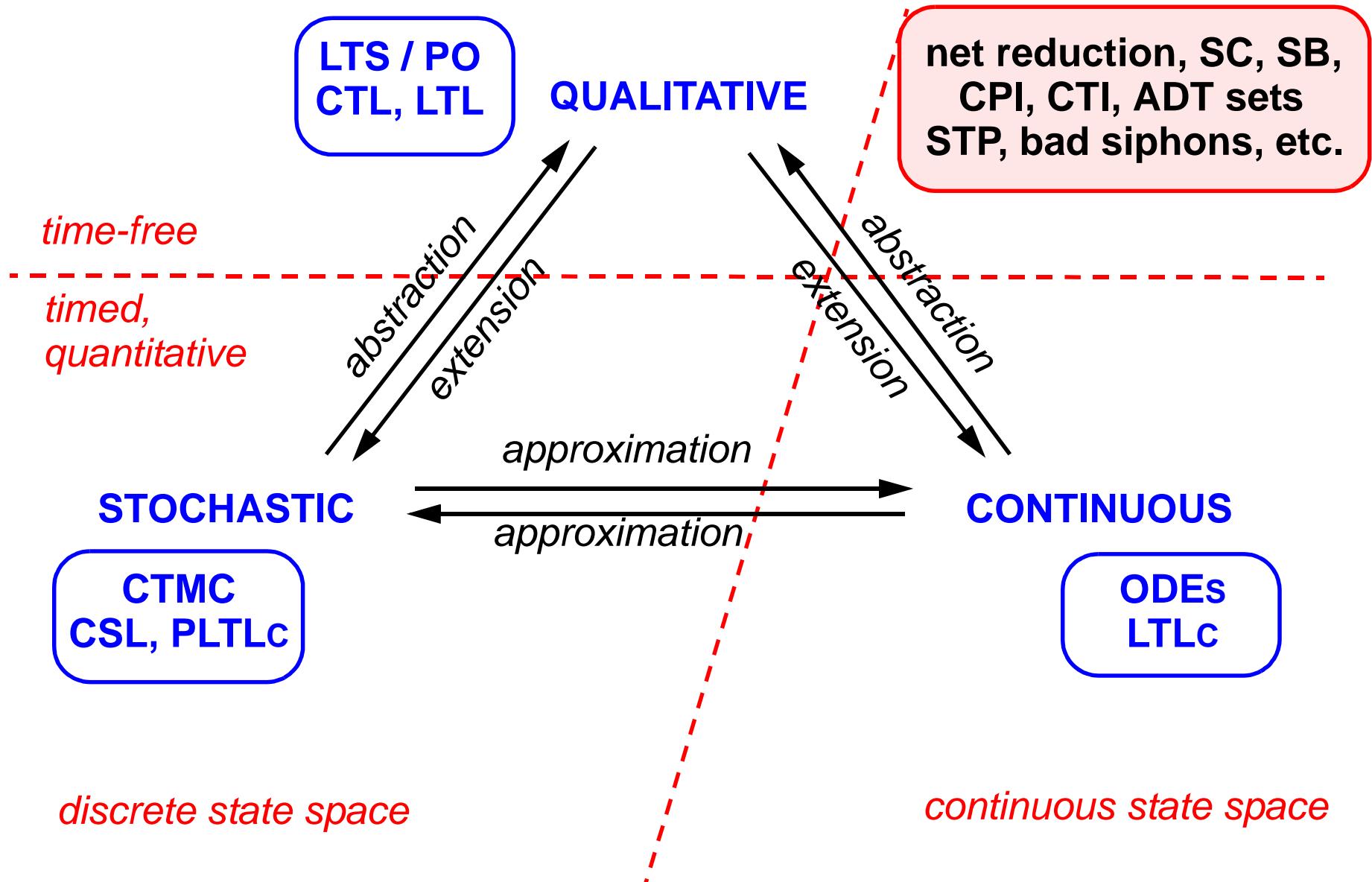
CONTINUOUS

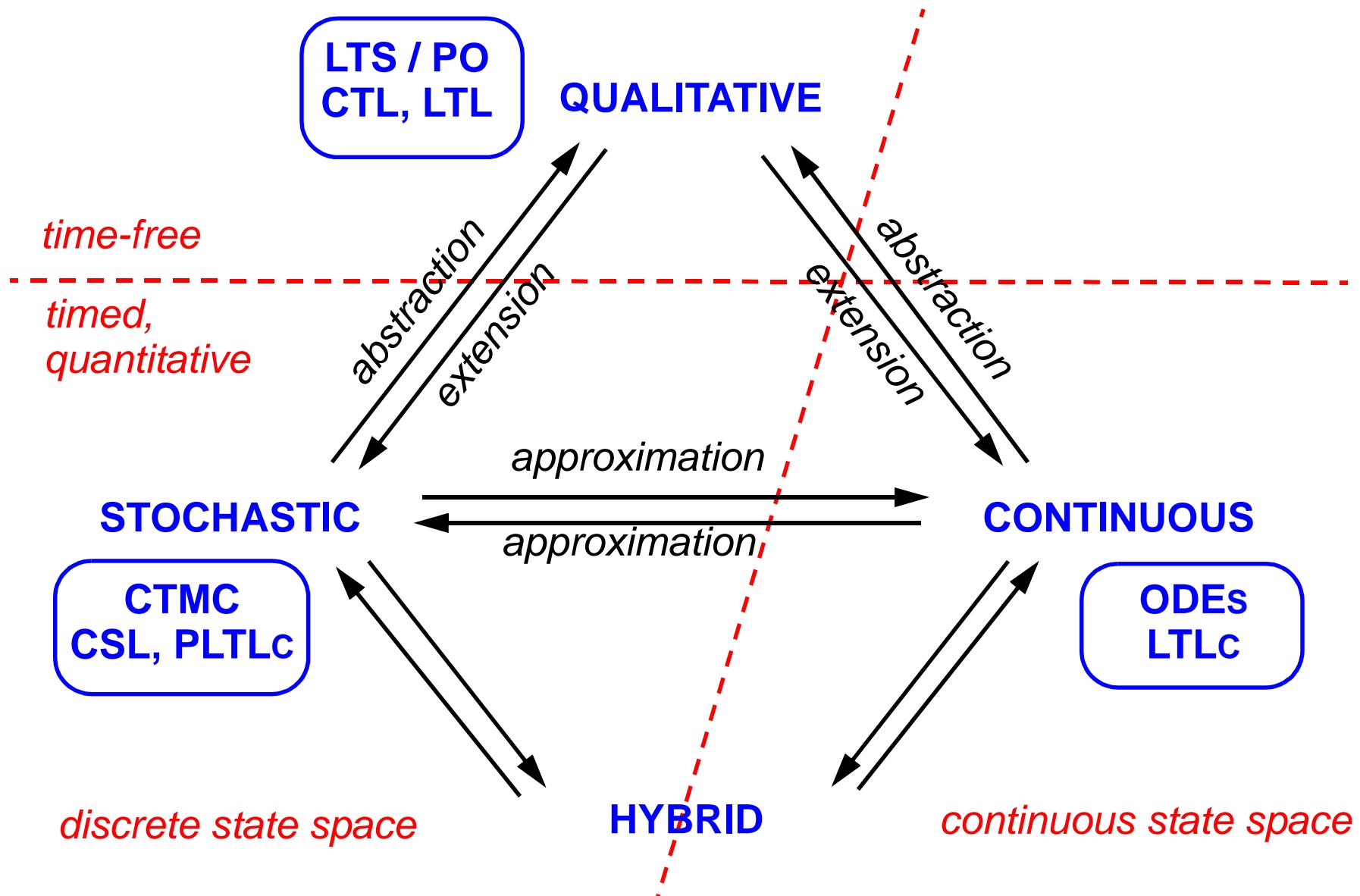


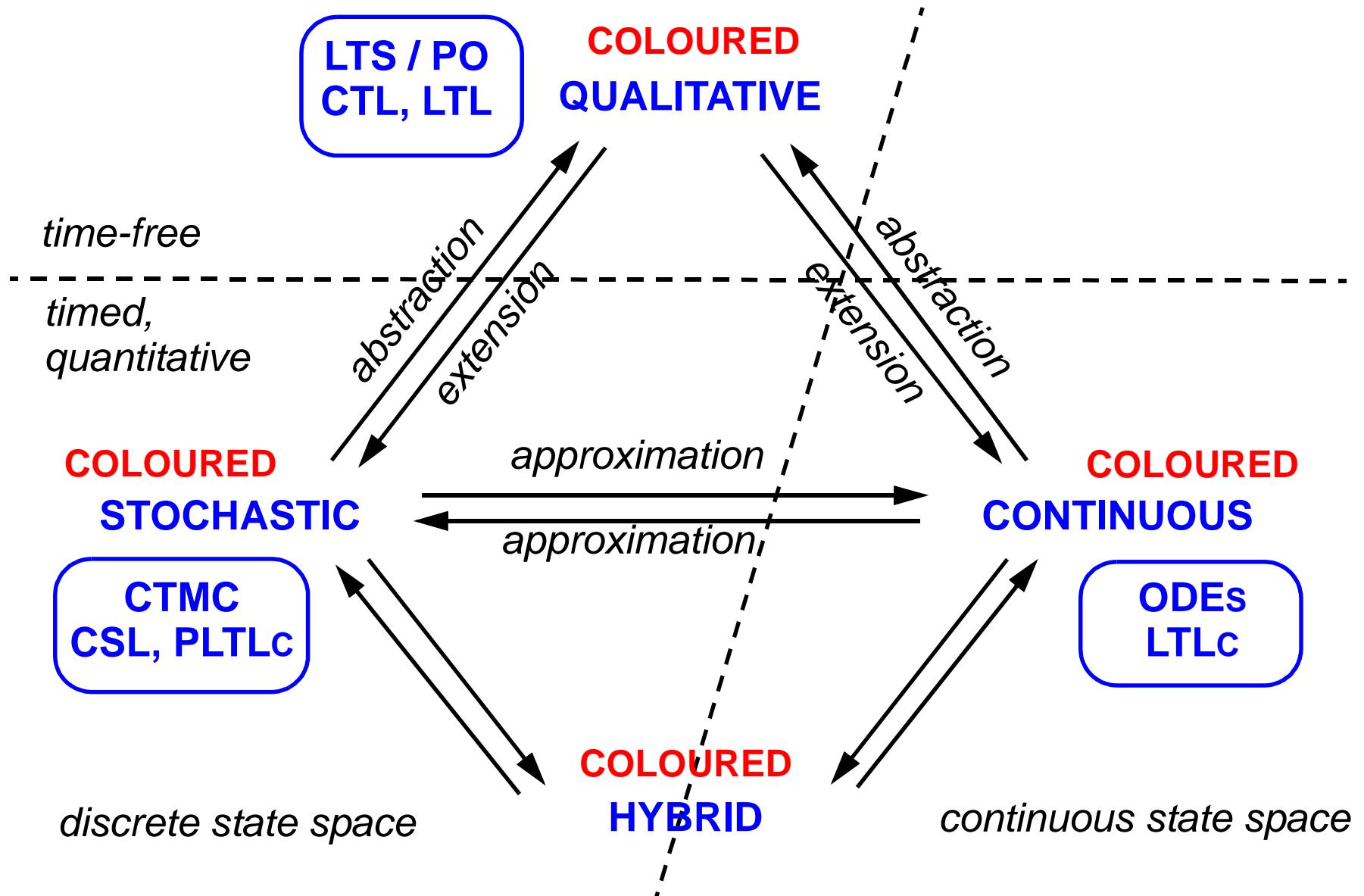






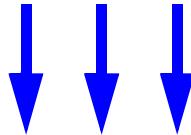






4X2

MODELS SHARING STRUCTURE



QUANTITATIVE MODEL = QUALITATIVE MODEL

+

**RATE FUNCTIONS
(KINETICS)**

OUR TOOLBOX

□ SNOOPY

- > *modelling and animation/simulation of hierarchical graphs,*
e.g. various Petri net classes, e.g. PN, XPN, SPN, XSPN, CPN, ...

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□ S4

- > *standalone, computational steering server*

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

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- > *PN, XPN, SPN, XSPN, SRN*
- > *symbolic and simulative model checking*

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□ Patty

- > *animation via web browser*

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- S4

- > *standalone, computational steering server*

- CTALENT

- > *PN, XPN, Timed Petri nets, T-Nets*

- > *mostly standard analysis techniques of Petri net theory*

- MARCIE

- > *PN, XPN, SPN, XSPN, SRN*

- > *symbolic and simulative model checking*

- Patty

- > *animation via web browser*

SBML import/export

EXPORT TO MATLAB AND

MANY OTHER TOOLS





- **readable & unambiguous**
-> *fault avoidant model construction*
- **locality - causality - concurrency**
- **compositional, hierarchical notations**
-> *logical and macro nodes*
- **executable**
-> *animation, simulation (token game)*



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- Petri net theory -> model validation**
 - > *P/T-invariants, partial order interpretation of T-invariants, conclusions CTI/CPI -> behavioural properties*
 - > *Siphon/Trap Property (STP), reduction rules, . . .*



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 - > *P/T-invariants, partial order interpretation of T-invariants, conclusions CTI/CPI -> behavioural properties*
 - > *Siphon/Trap Property (STP), reduction rules, . . .*
- **umbrella with unifying power**
 - > *interpretation in qualitative / stochastic / continuous / hybrid paradigm*

T- INVARIANTS

(ELEMENTARY MODES)

(EXTREME PATHWAYS)

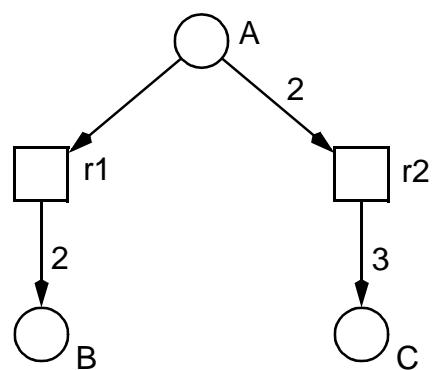
(GENERIC PATHWAYS)

$r1: A \rightarrow 2 B$

$r2: 2 A \rightarrow 3 C$

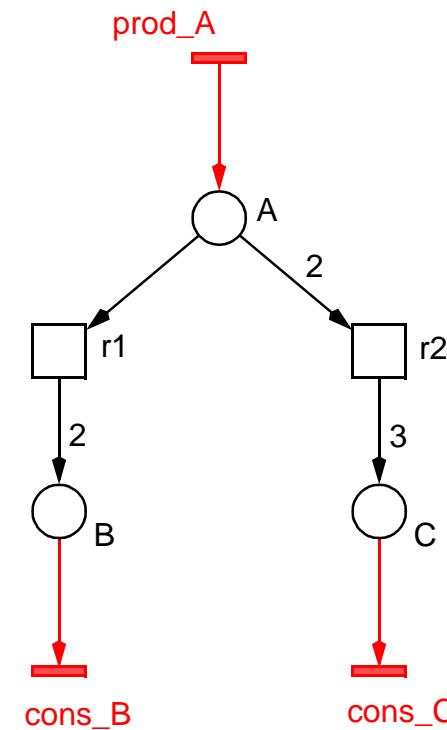
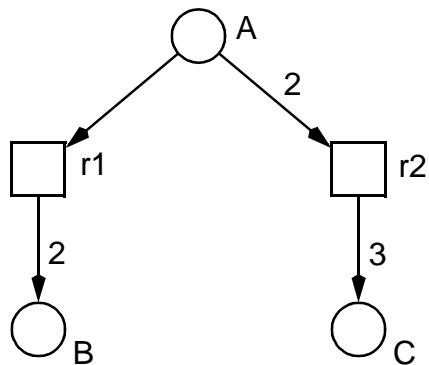
$r1: A \rightarrow 2 B$

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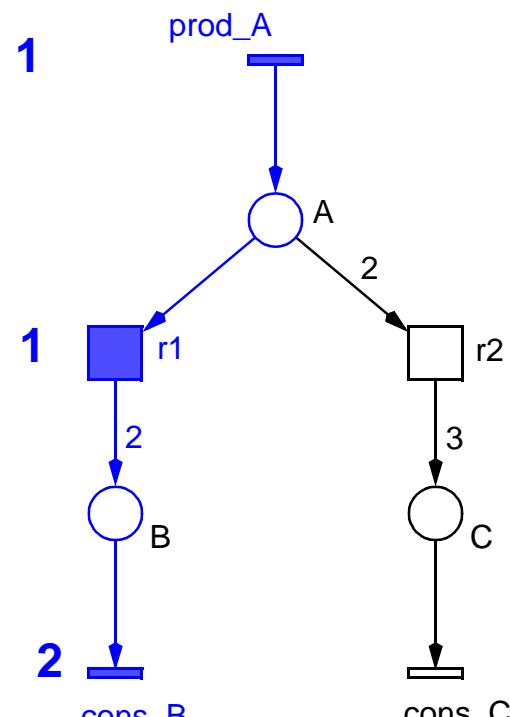
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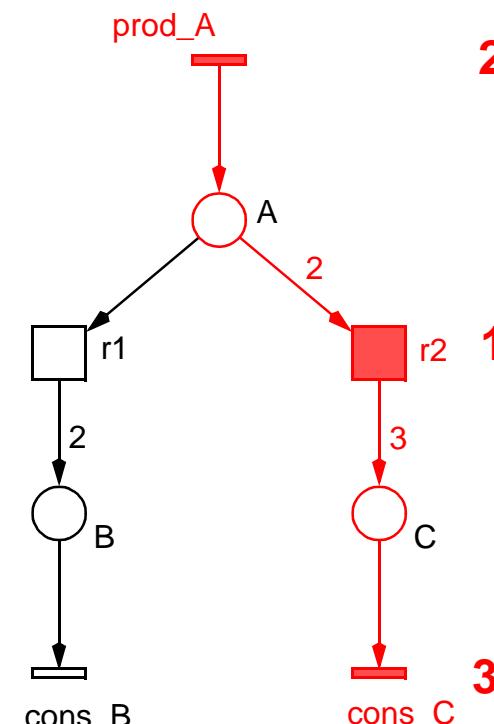


$r1: A \rightarrow 2 B$

$r2: 2 A \rightarrow 3 C$



T-INVARIANT 1



T-INVARIANT 2

INCIDENCE MATRIX C

PN & BioModel Engineering

- a representation of the net structure

=> stoichiometric matrix

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

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

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

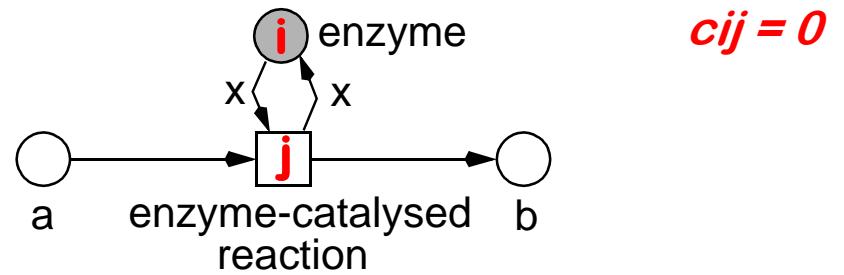
- matrix entry c_{ij} :

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



- Lautenbach, 1973 -> Schuster, 1993
 - T-invariant x -> *multiset of transitions*
 - > integer solution of $Cx = 0, x \neq 0, x \geq 0$
 - support of a T-invariant x -> $\text{supp}(x)$ -> *set of transitions*
 - > set of transitions involved, i.e. $x(i) \neq 0$
 - minimal T-invariants
 - > there is no T-invariant with a smaller support
 - > 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
$$kx = \sum_i a_i x_i$$

□ T-invariants = (multi-) sets of transitions = Parikh vector

- > zero effect on marking
- > reproducing a marking / system state

□ two interpretations

1. *partially ordered transition sequence* -> behaviour understanding
of transitions occurring one after the other
-> substance / signal flow
2. *relative transition firing rates* -> steady state behaviour
of transitions occurring permanently & concurrently
-> steady state behaviour

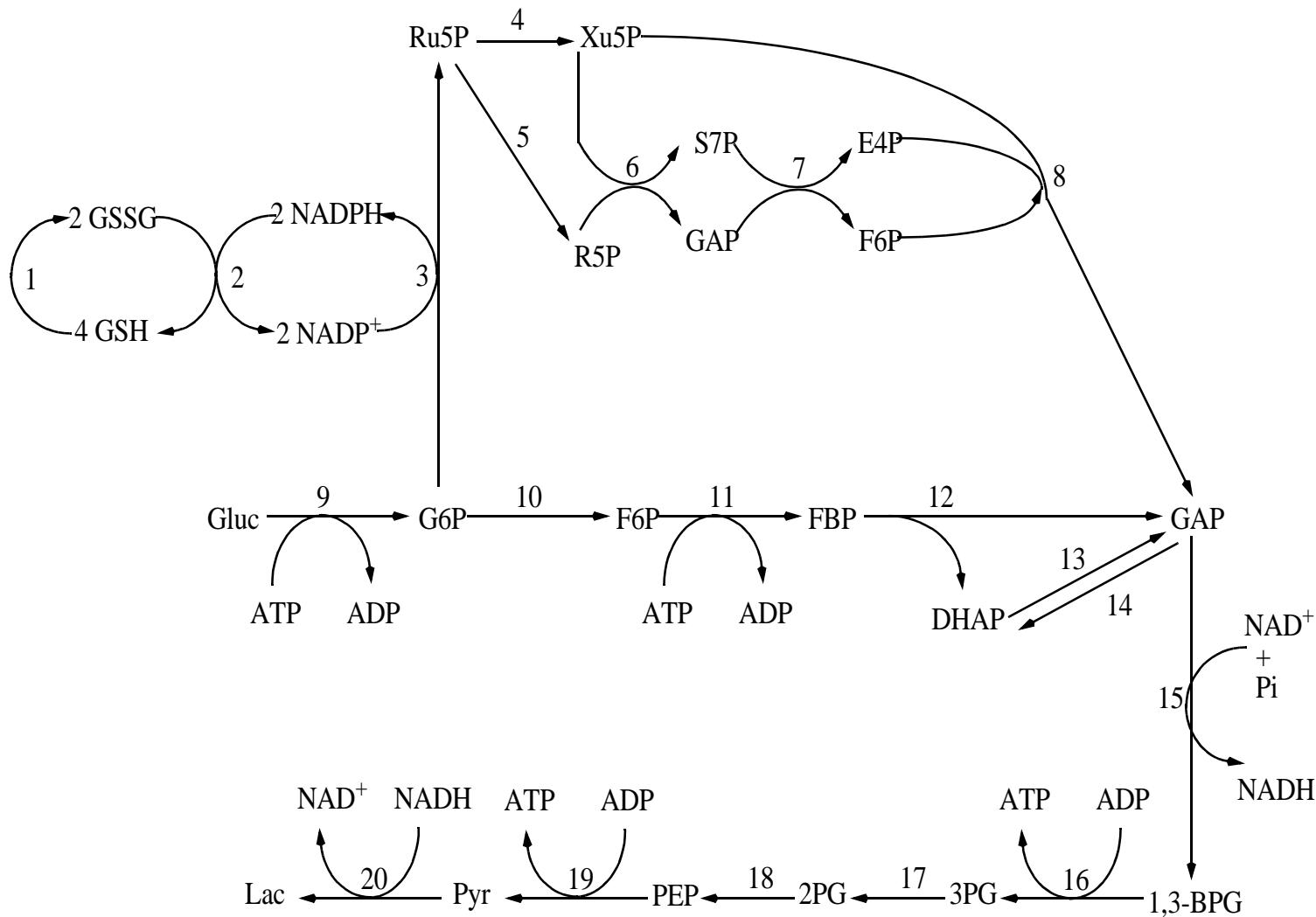
□ a minimal T-invariant defines a connected subnet

- > 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

Ex1 - Glycolysis and Pentose Phosphate Pathway

PN & BioModel Engineering

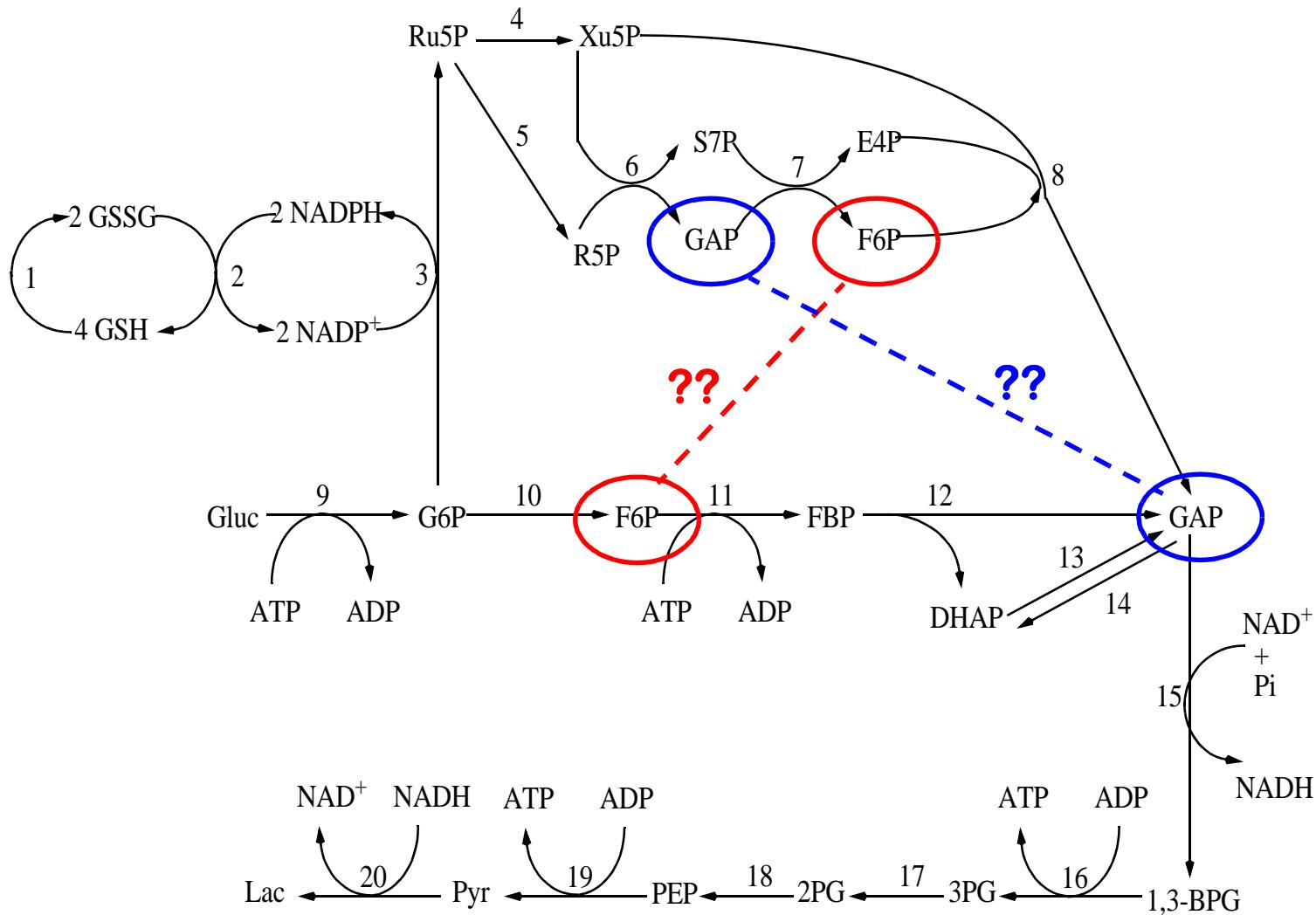
[Reddy 1993]



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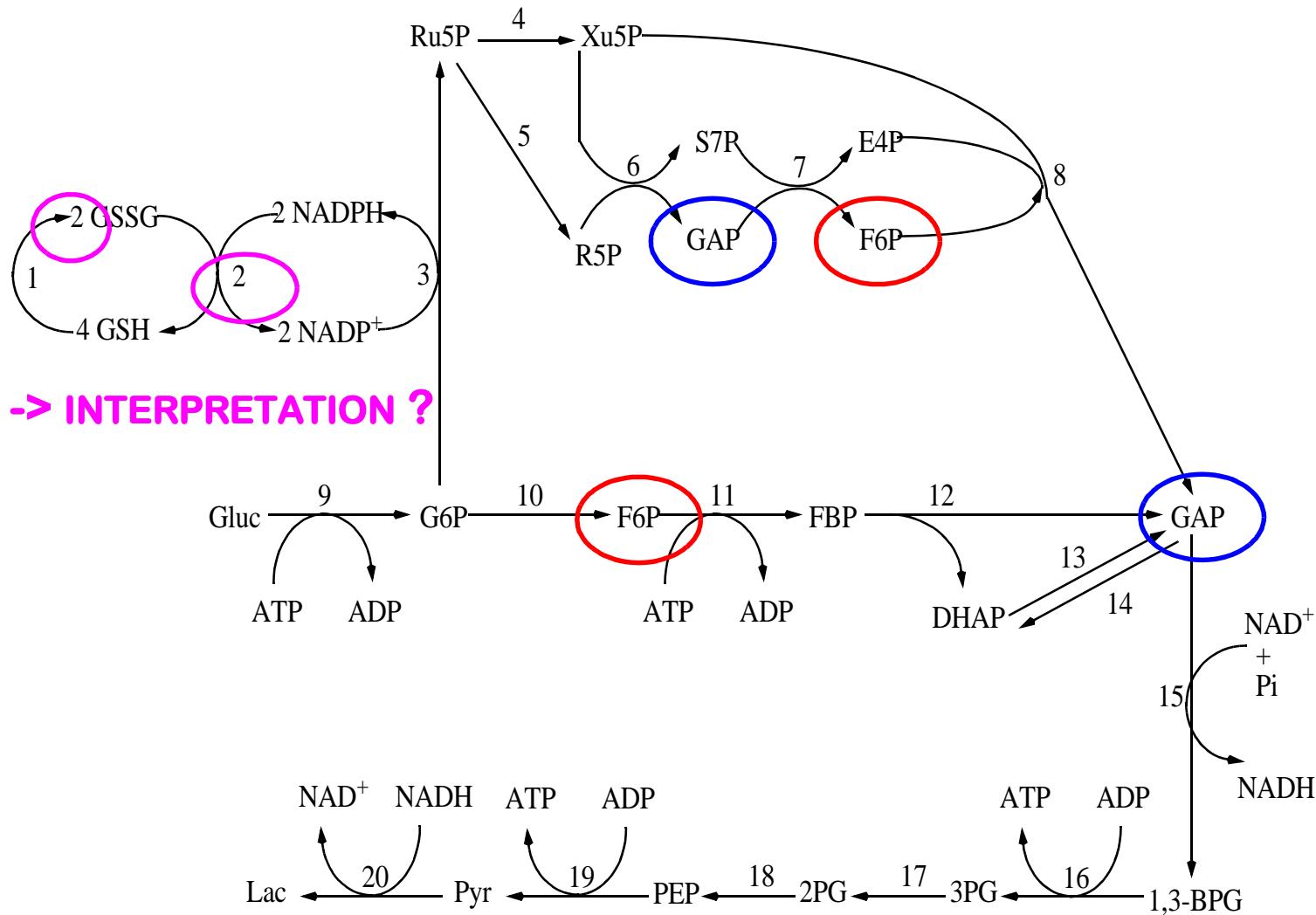
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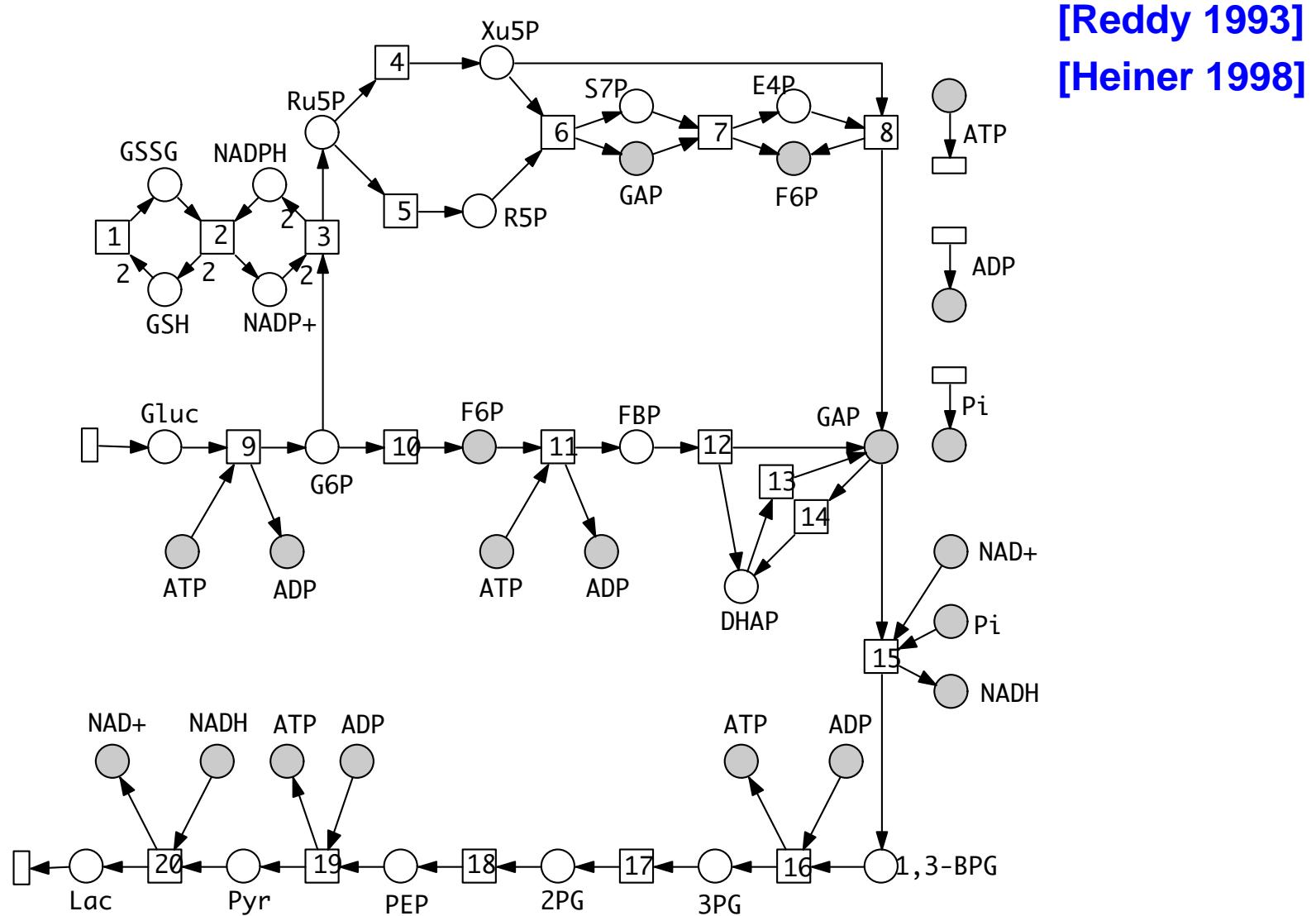
PN & BioModel Engineering

[Reddy 1993]



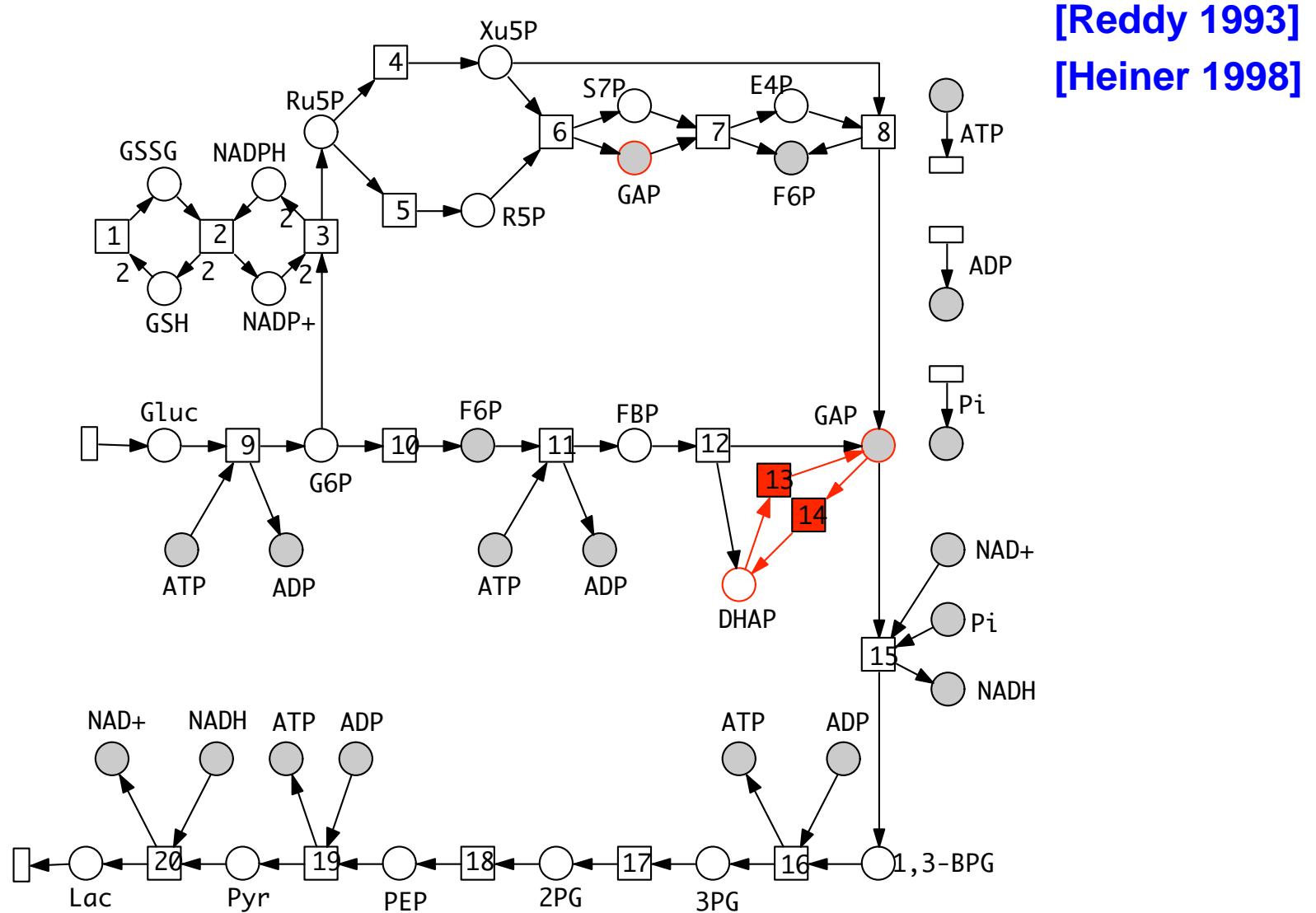
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PN & BioModel Engineering



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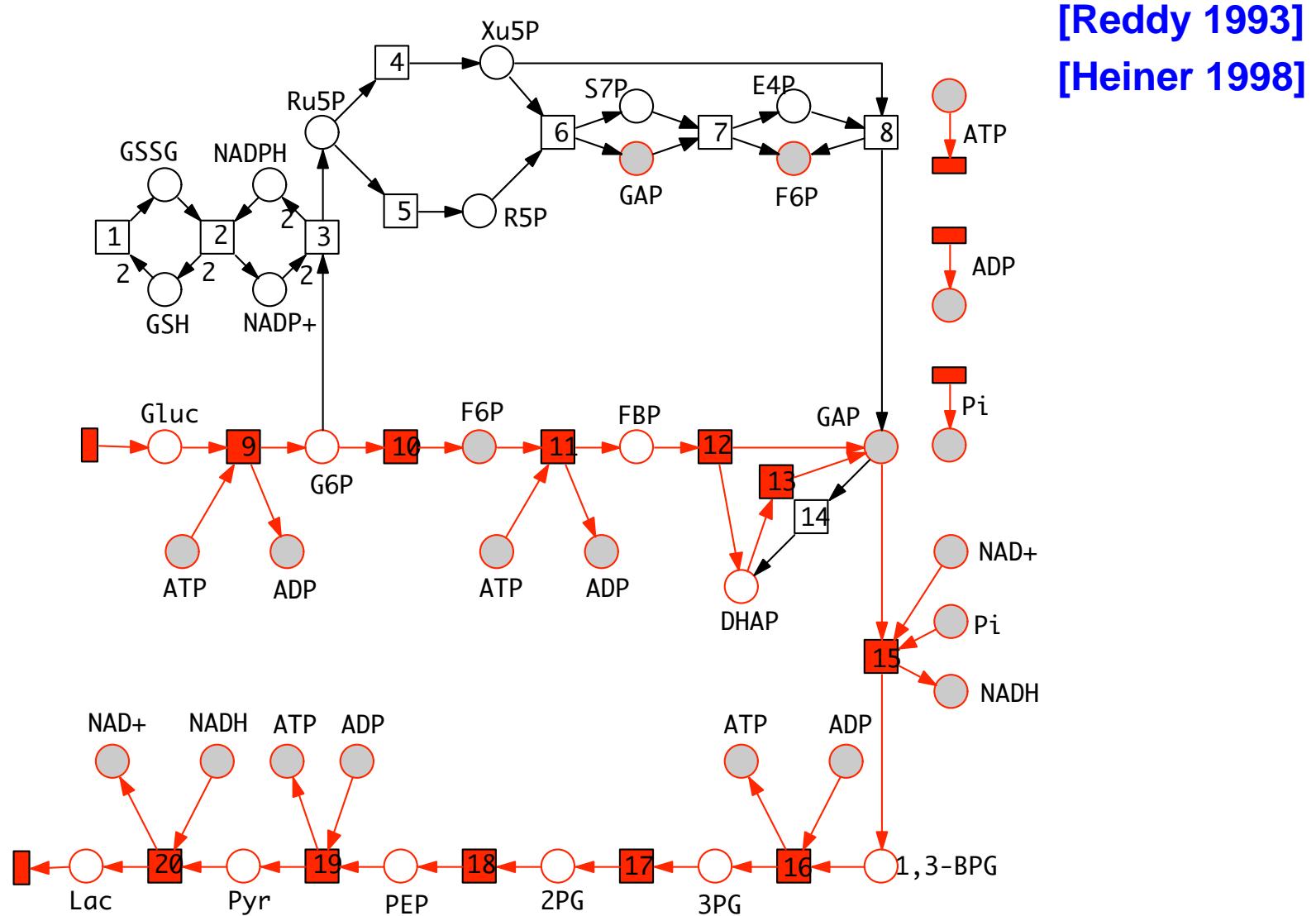


[Reddy 1993]

[Heiner 1998]

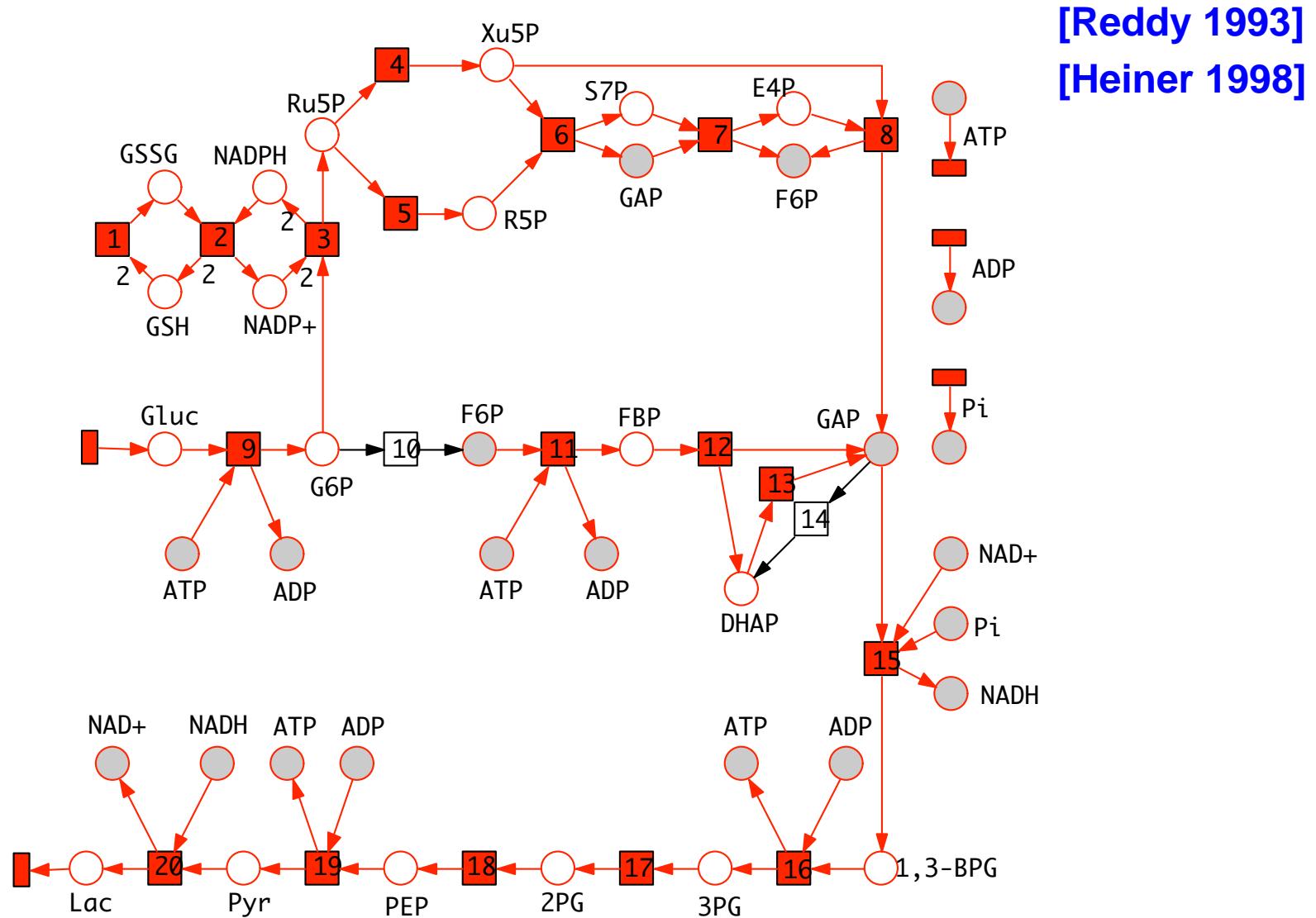
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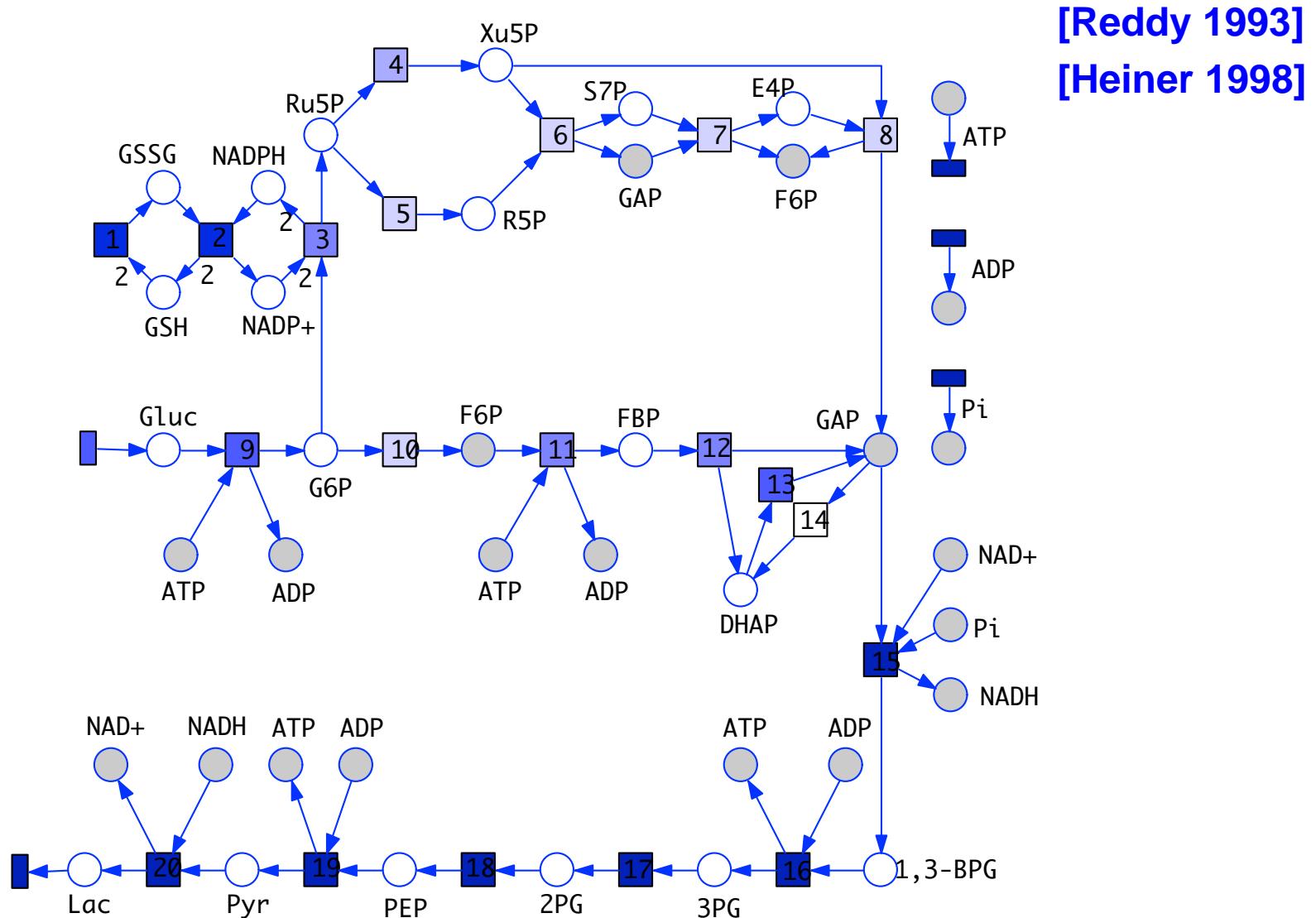
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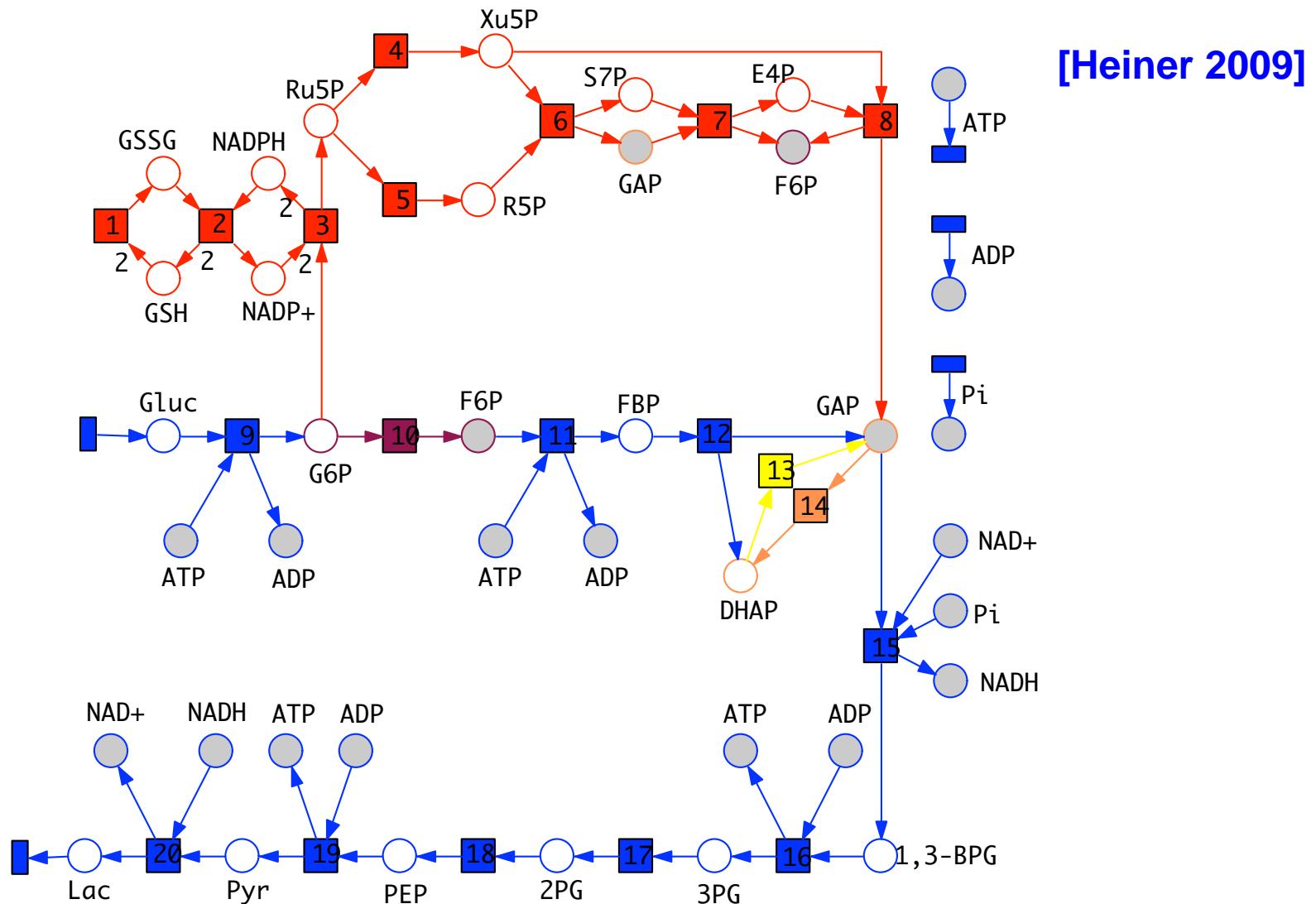
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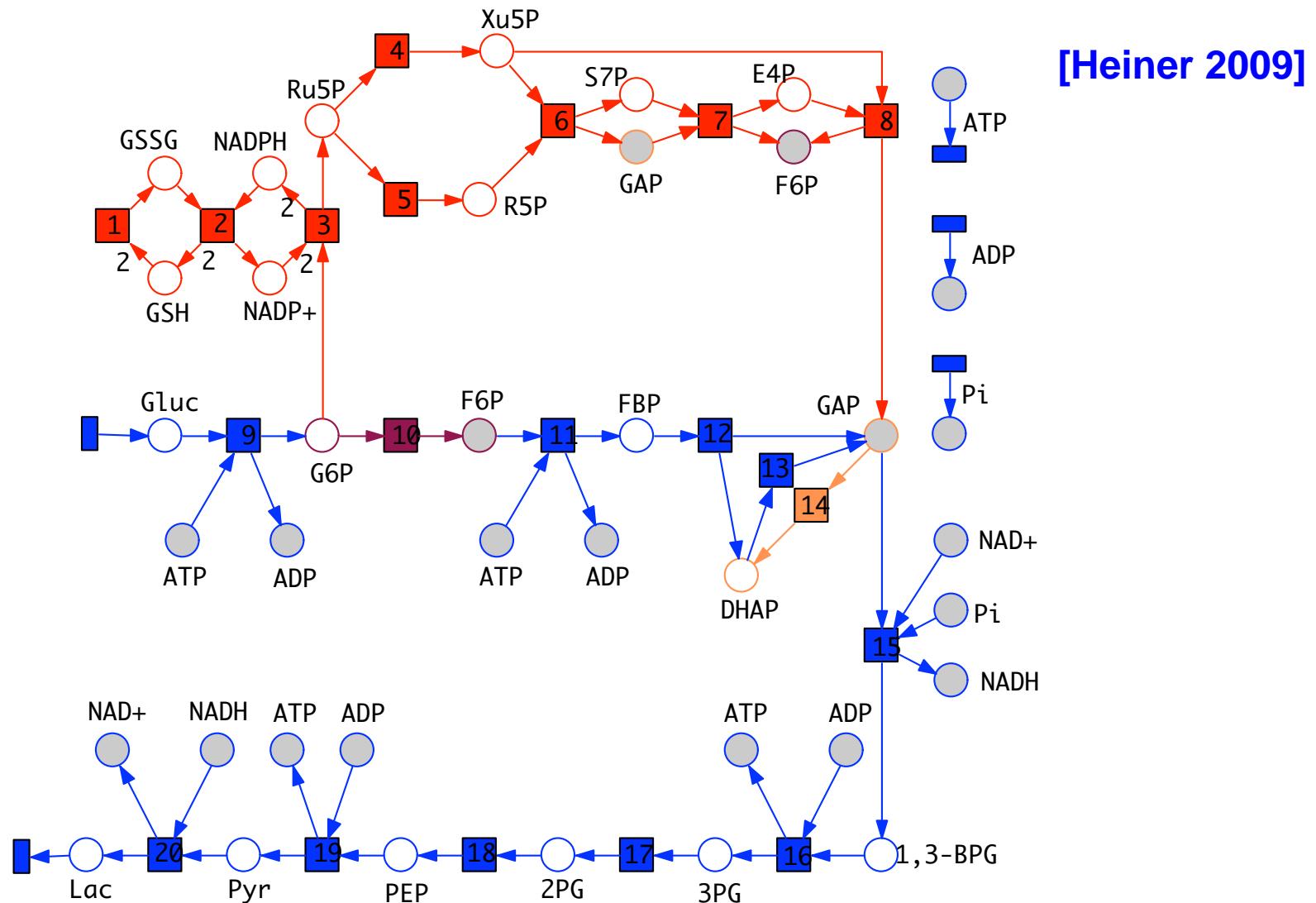
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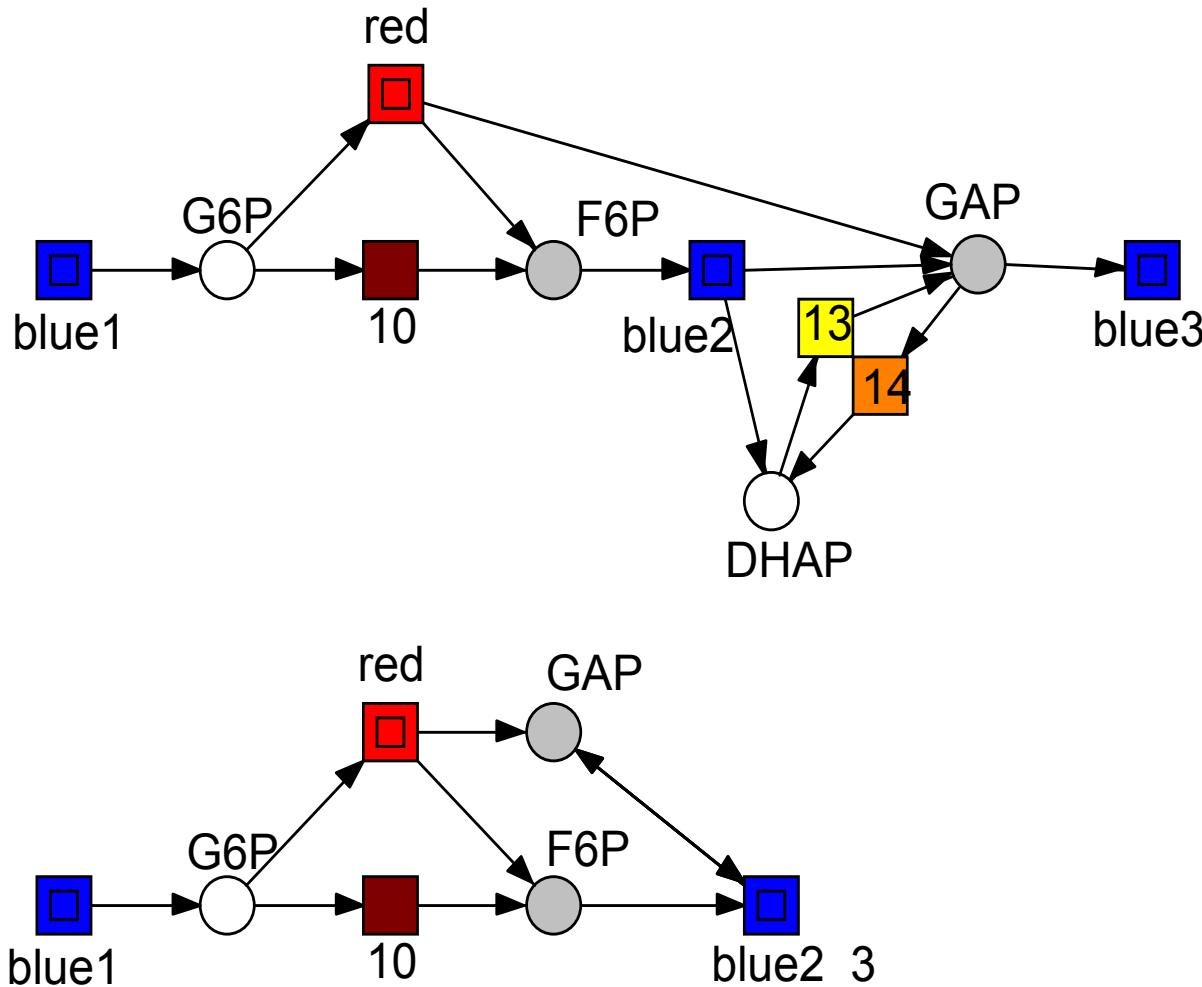
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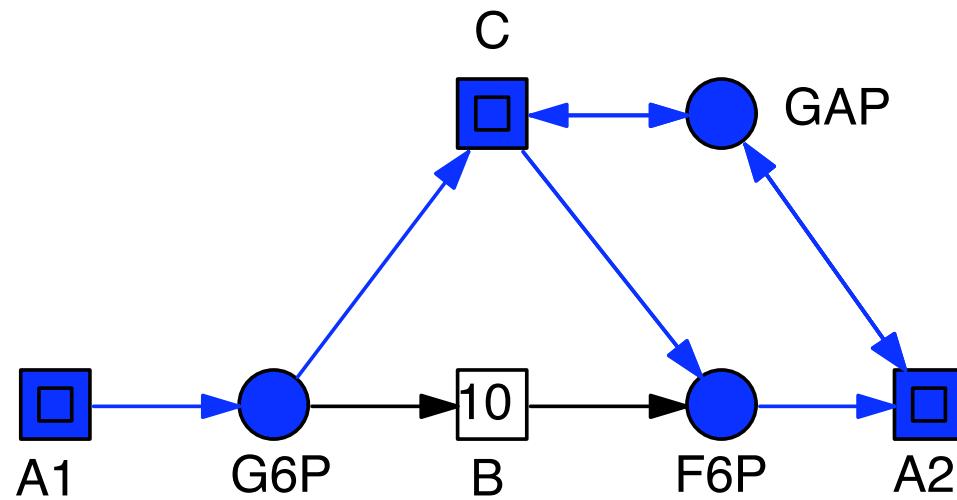
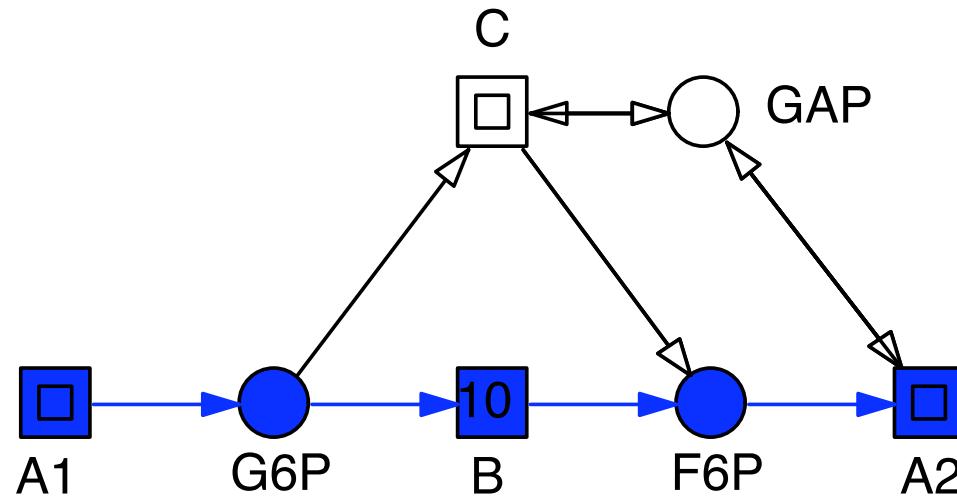
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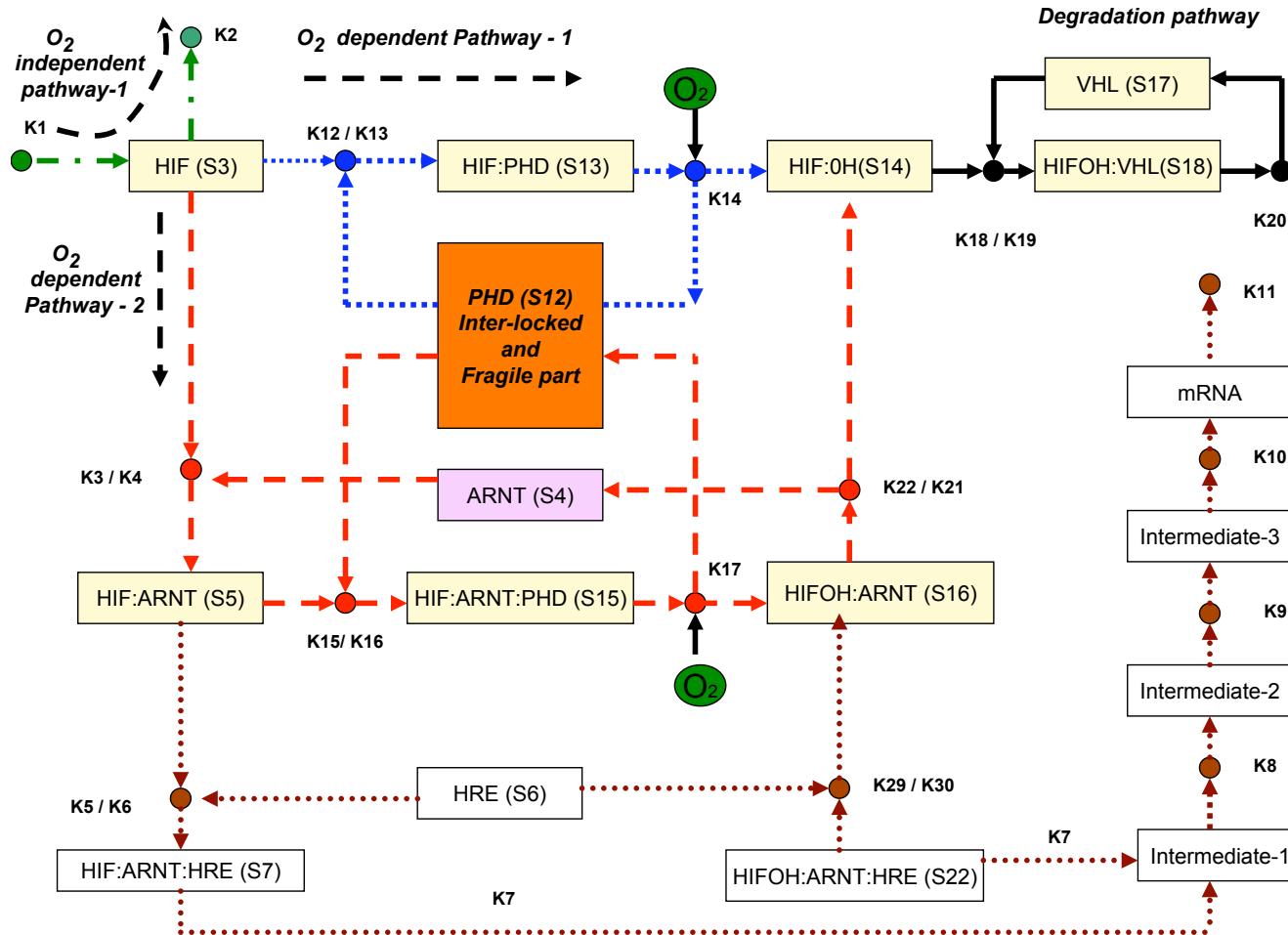
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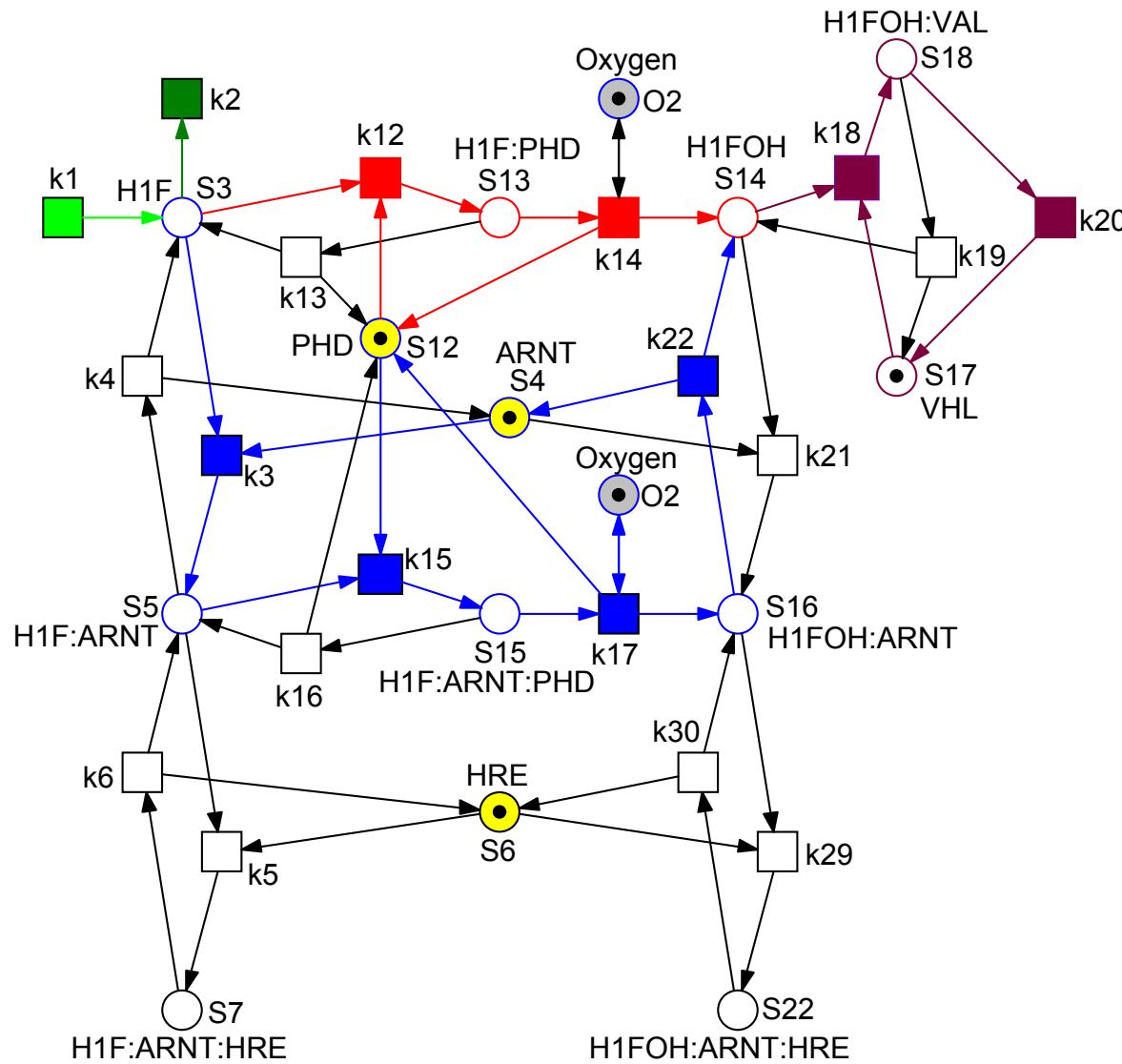


ABOUT THE RELATION QUALITATIVE VS CONTINUOUS

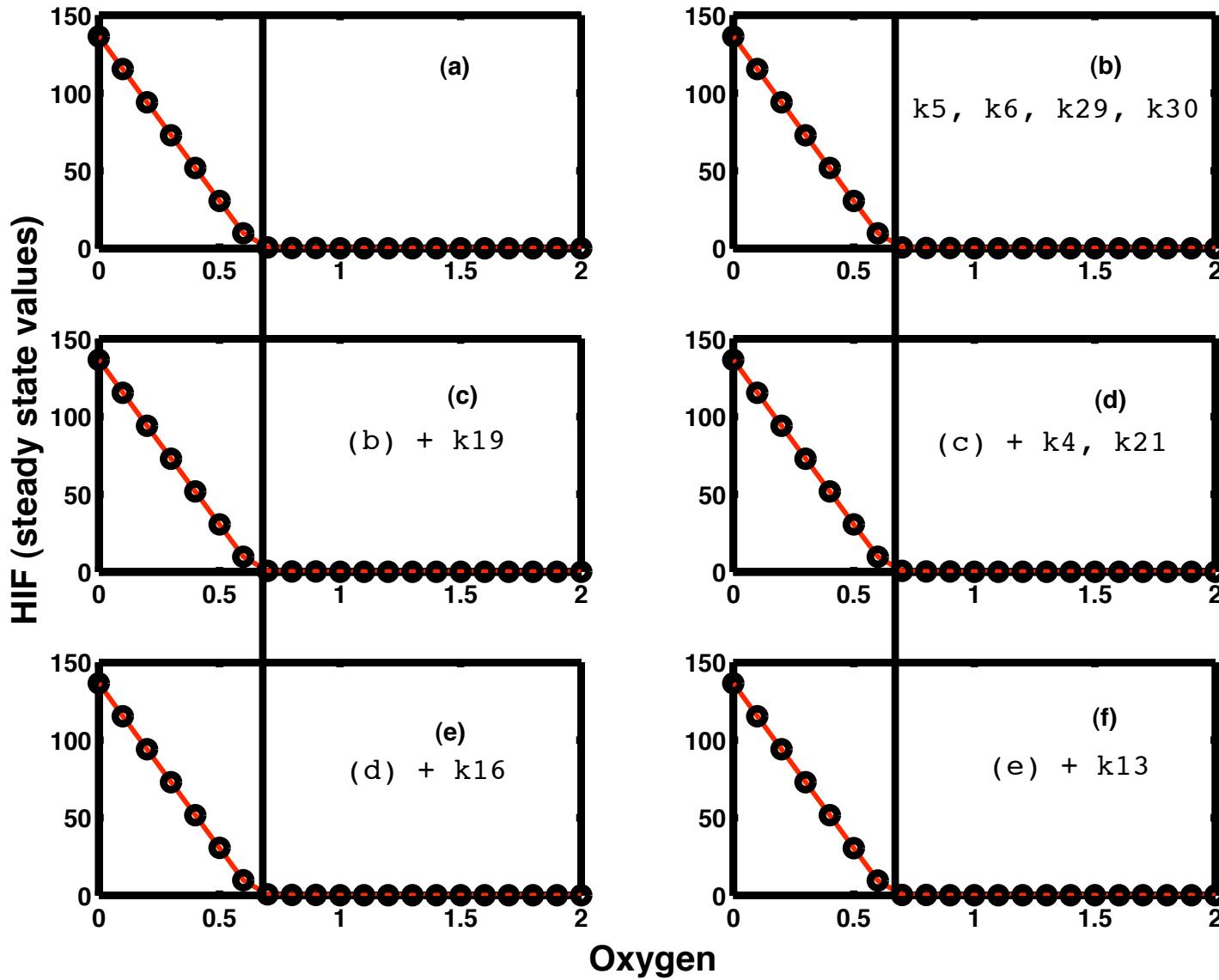
[YU ET AL. 2007]



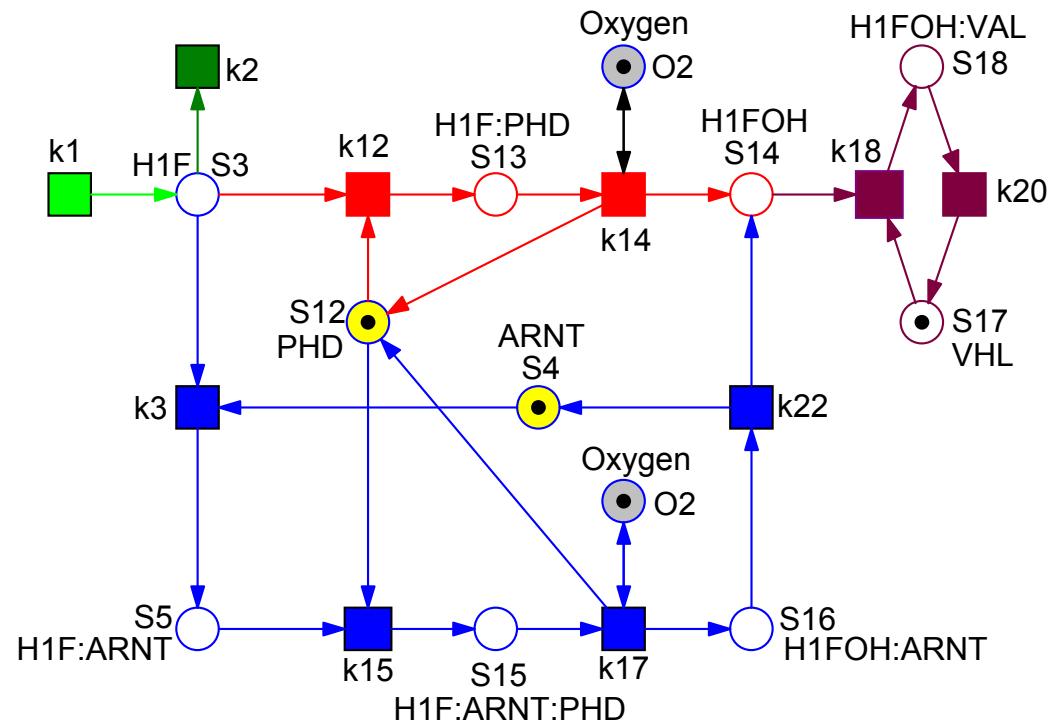
[HEINER,
SRIRAM 2010]



Ex4 - HYPOXIA

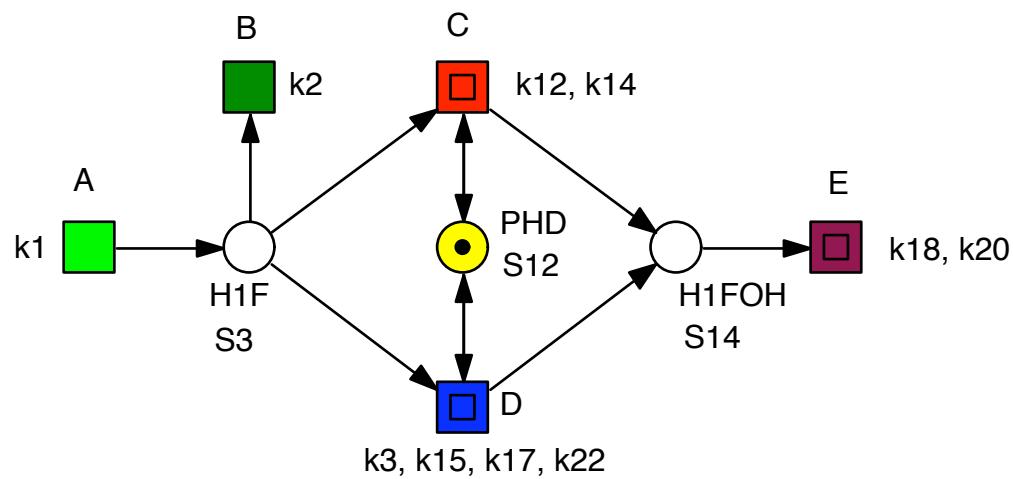


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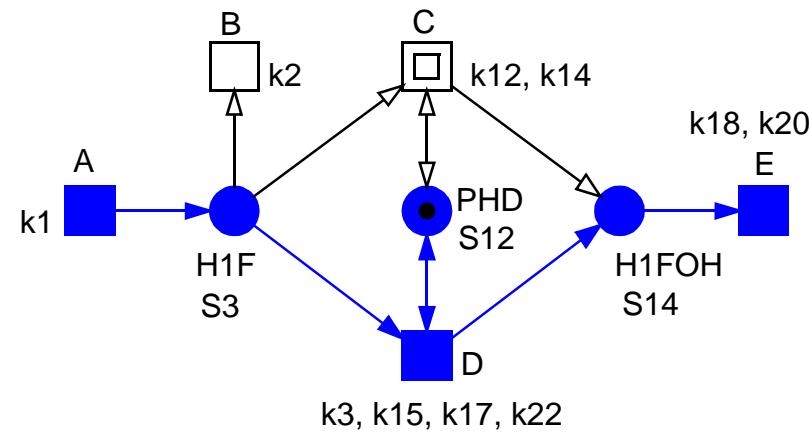
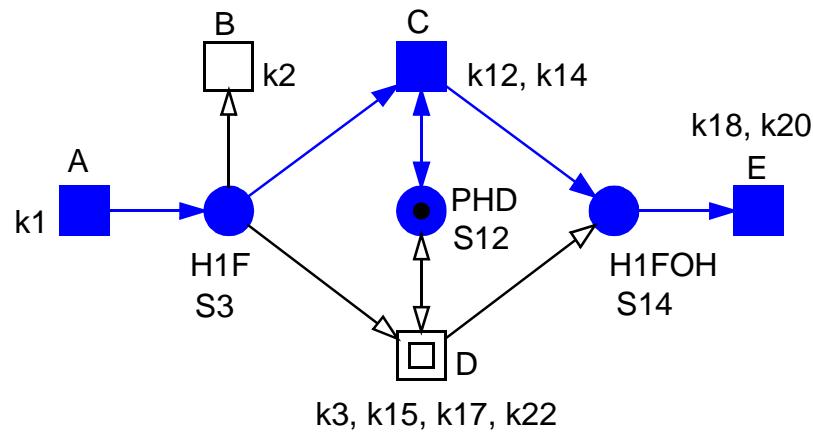
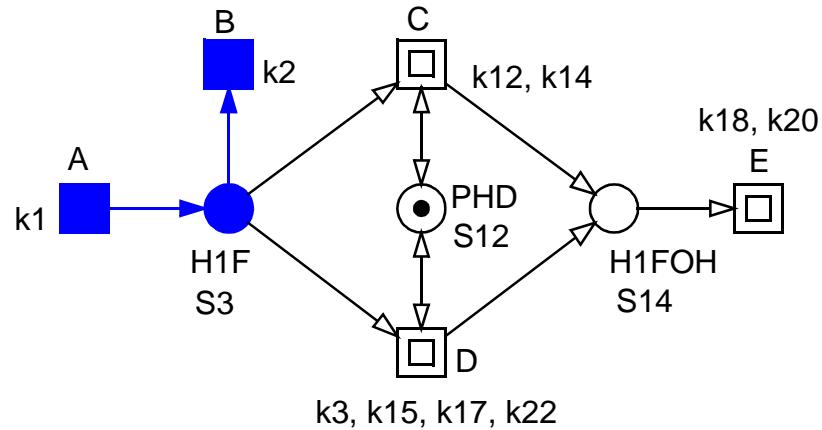
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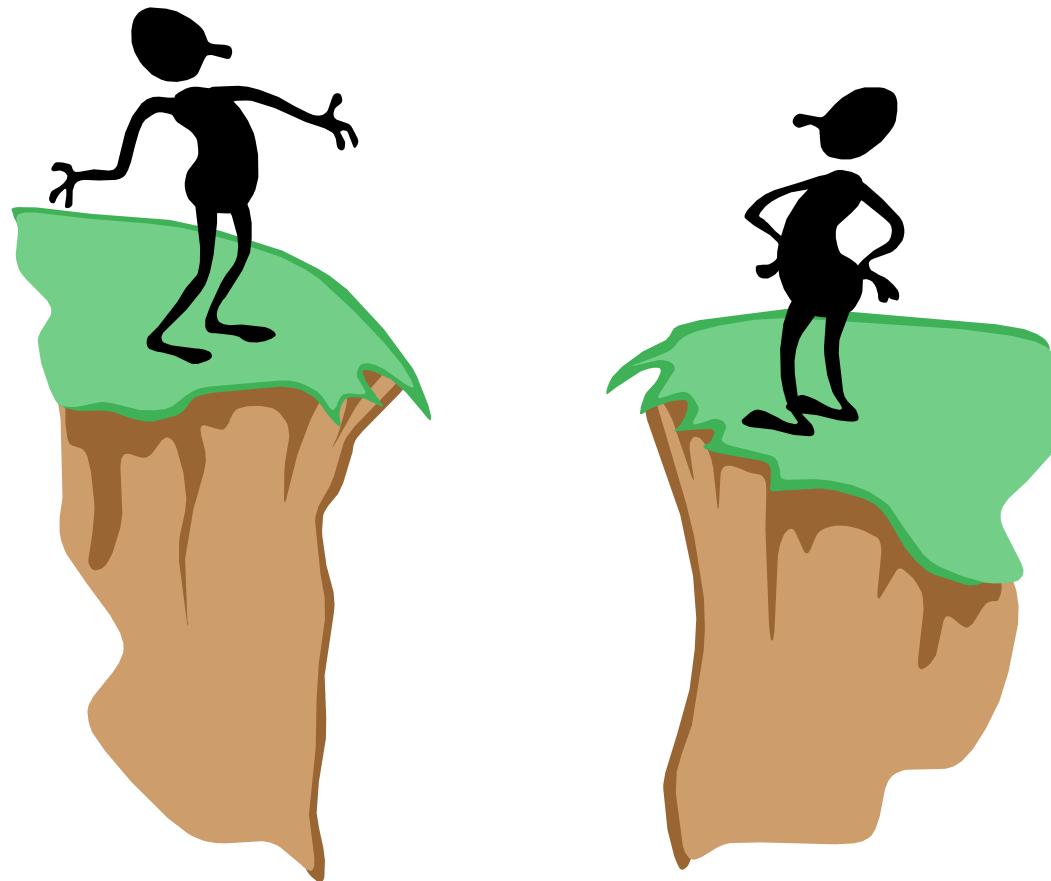
PN & BioModel Engineering



Ex4 - HYPOXIA

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THANKS !

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