

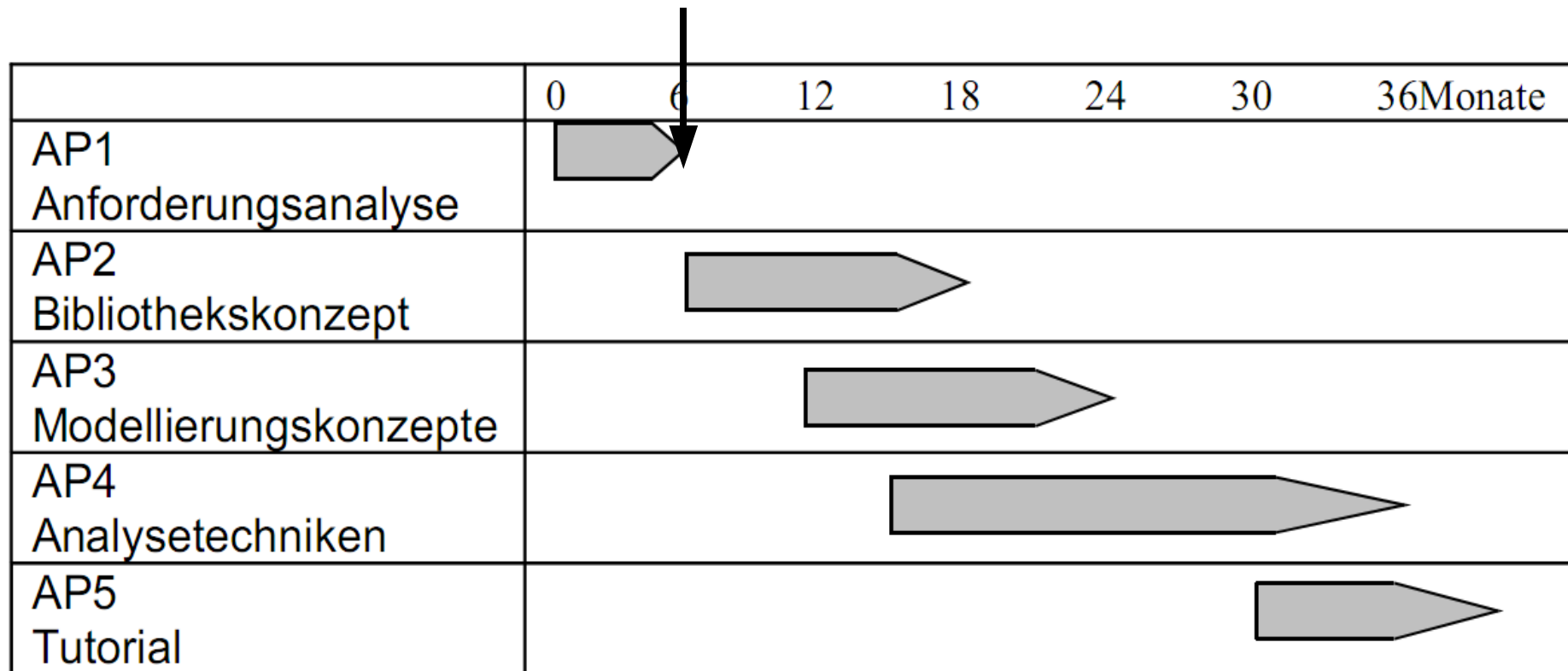
A Framework for Modular Modeling and Analysis of Signaling Networks

Cottbus
17 August 2009

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Project Milestones

- WP1 - Requirements analysis
- WP2 - Library approach for generic model components
- WP3 - Modeling concepts for dealing with model alternatives
- WP4 - Analysis techniques for identification and
Behavior comparison of model components
- WP5 - Tutorial



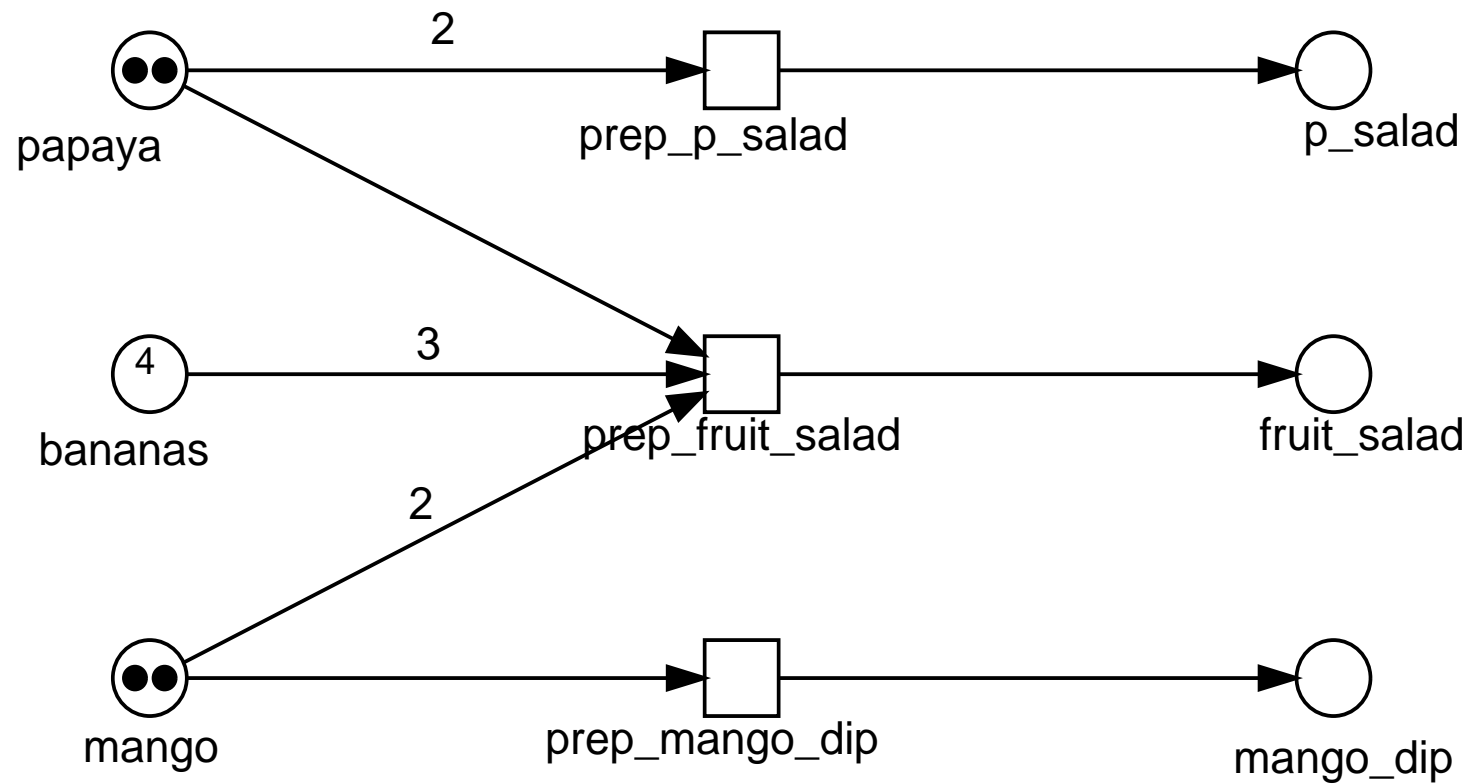
- Building a collection of the relevant network examples which are fully documented in the literature;
- Literature search for techniques to identify structure-based modules; evaluation on automatability, tool availability, and expense of implementation;
- Identification of structural components, manually or using tools, if possible;
- Identification of classes of structural components, which can be grouped together as generic model components;
- Develop a concept for the coupling with the database developed in M2;

Project Results

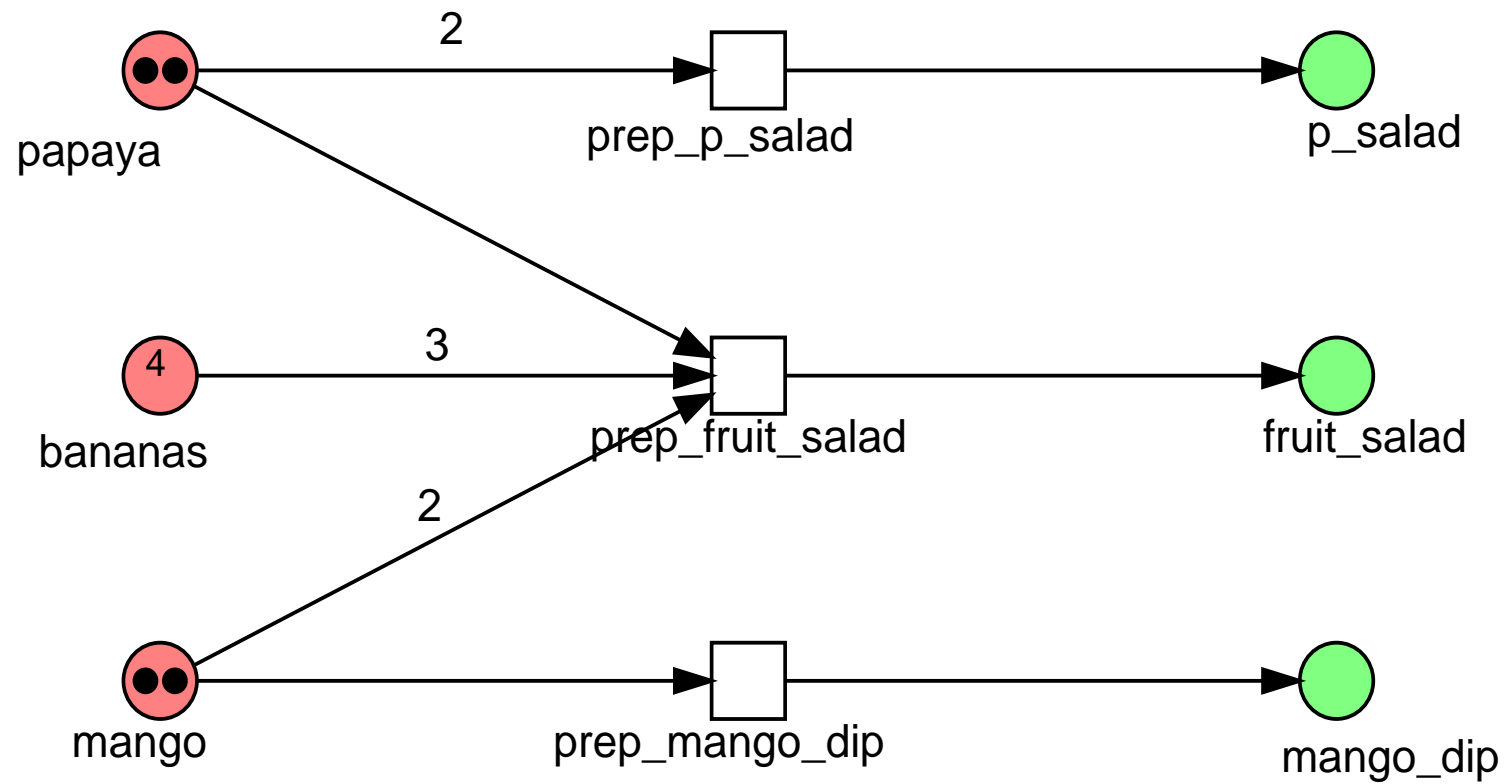
June 2009 --- August 2009

- build a more compact and parametric representation of a system by folding similar subnets
- it is possible to represent very concisely systems that would have required a huge uncoloured net
- these similar subnets when folded are distinguished by coloured tokens

→ Prepare salad

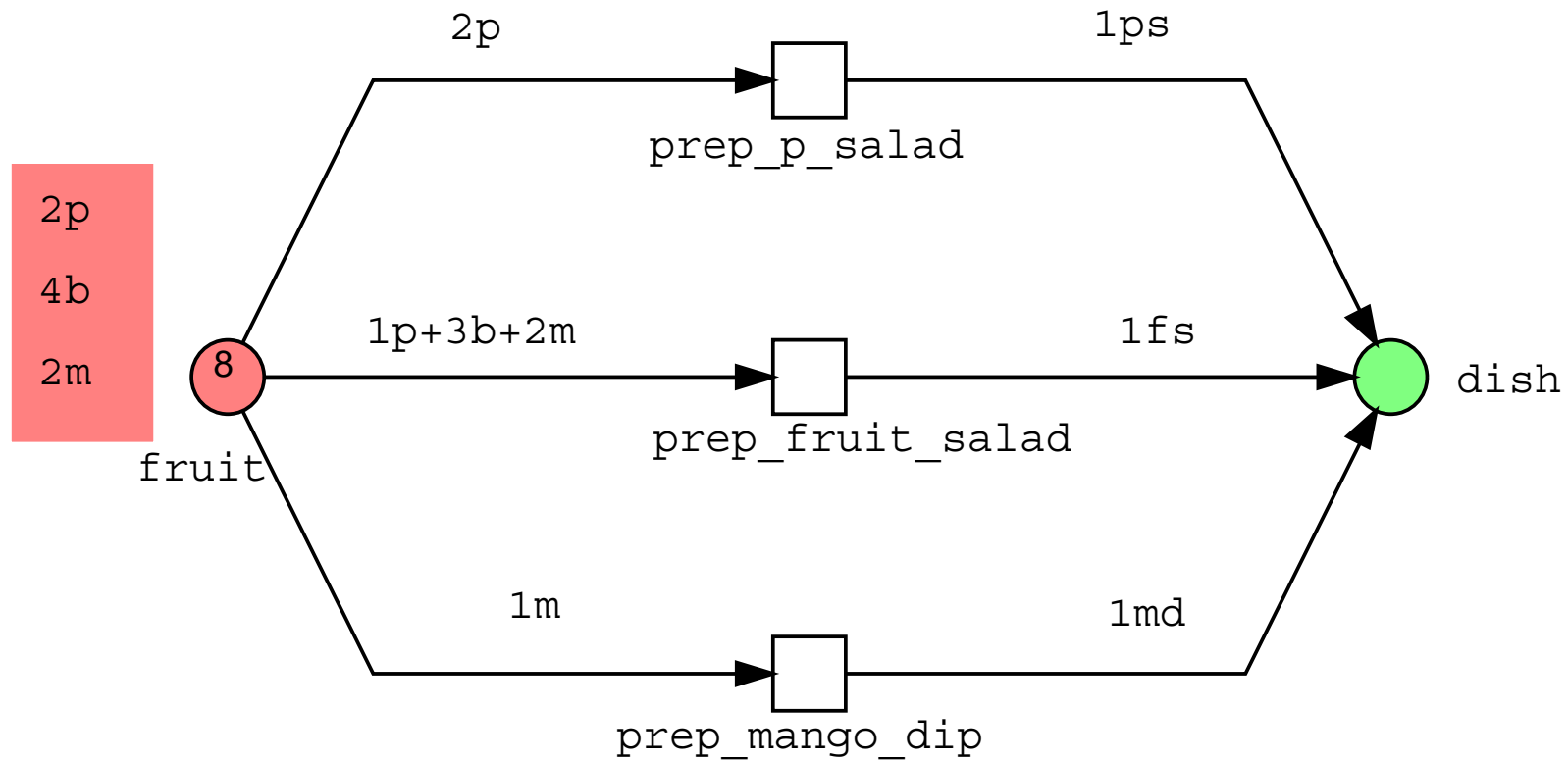


→ Prepare salad



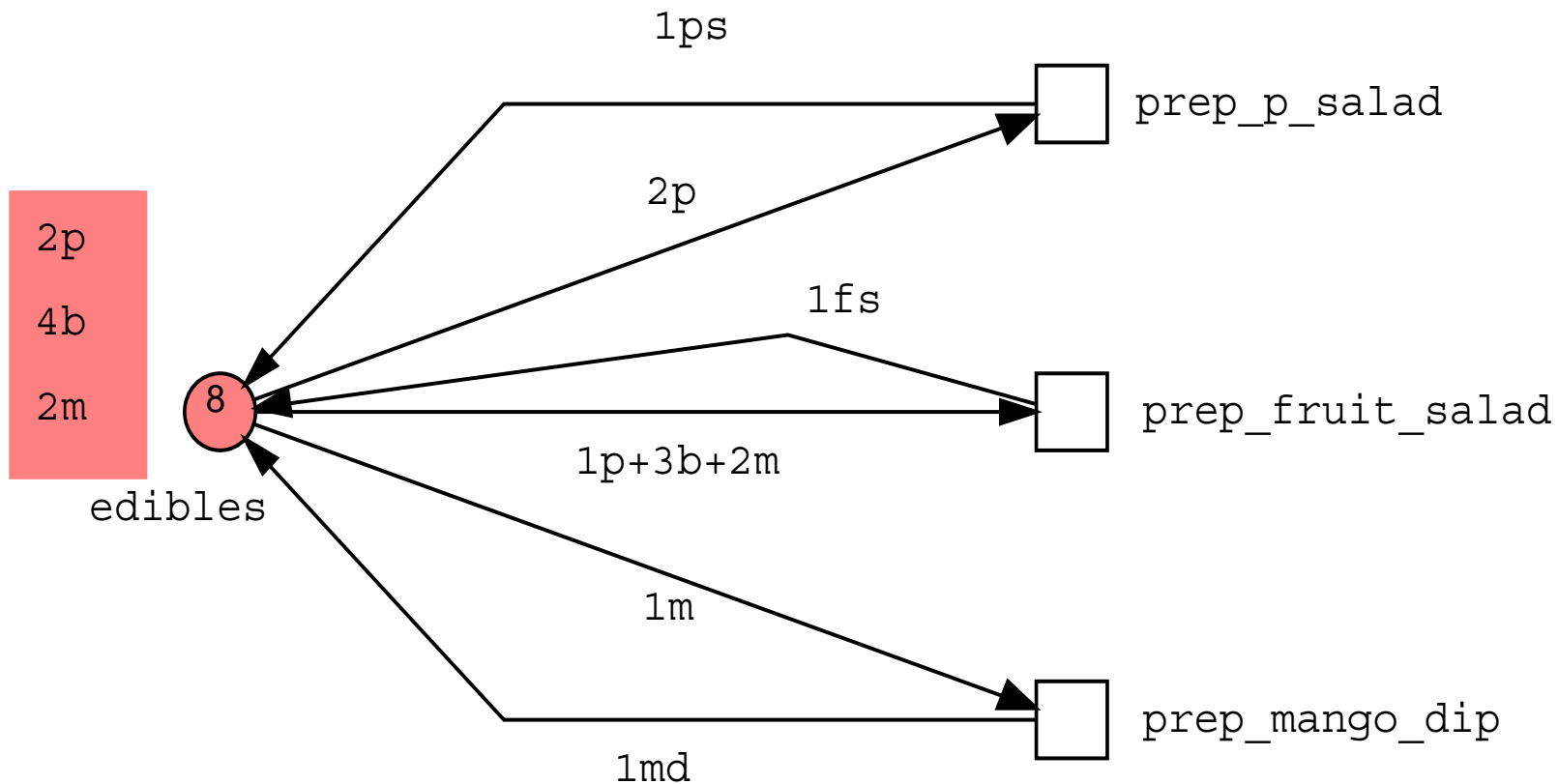
→ Prepare salad

Colourset:
 Enumeration fruit = {p, b, m}
 Enumeration dish = {ps, fs, md}



→ Prepare salad

Colourset:
 Enumeration fruit = {p, b, m}
 Enumeration dish = {ps, fs, md}
 union edibles = fruit, dish



→ Prepare salad

Colourset:

enumeration fruit = {p, b, m}

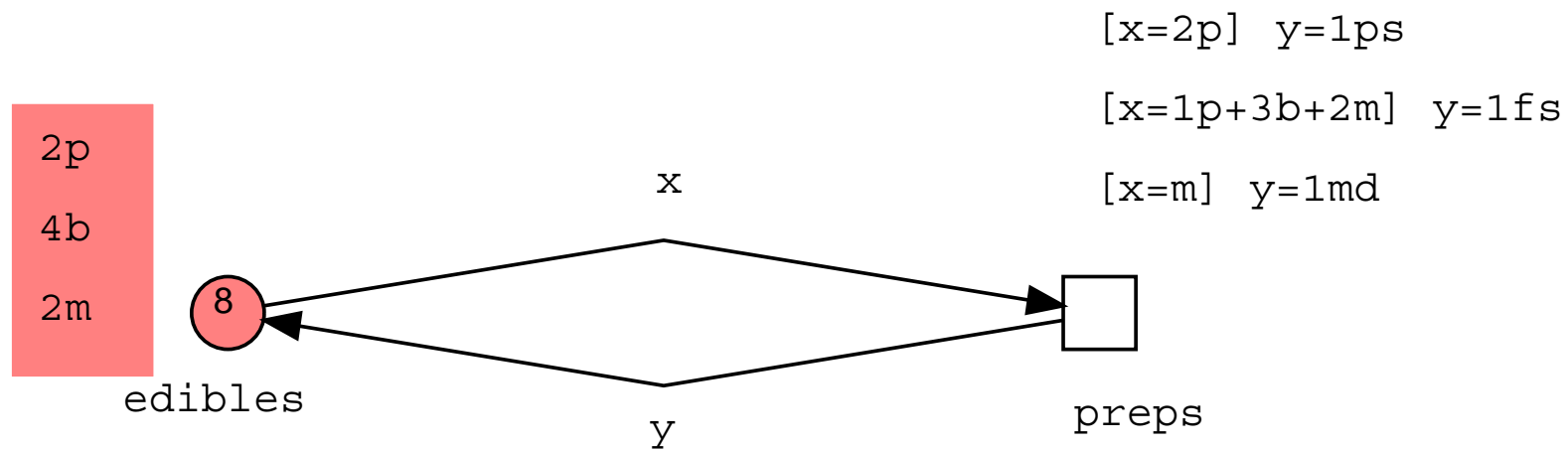
enumeration dish = {ps, fs, md}

union edibles = fruit, dish

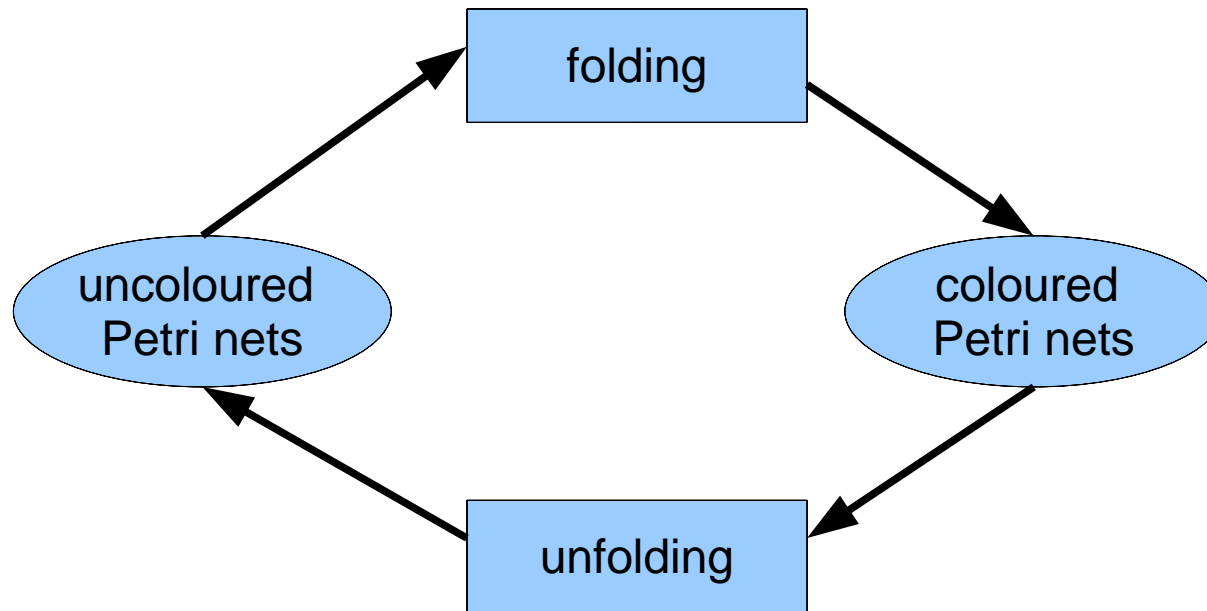
Variable:

fruit x

dish y

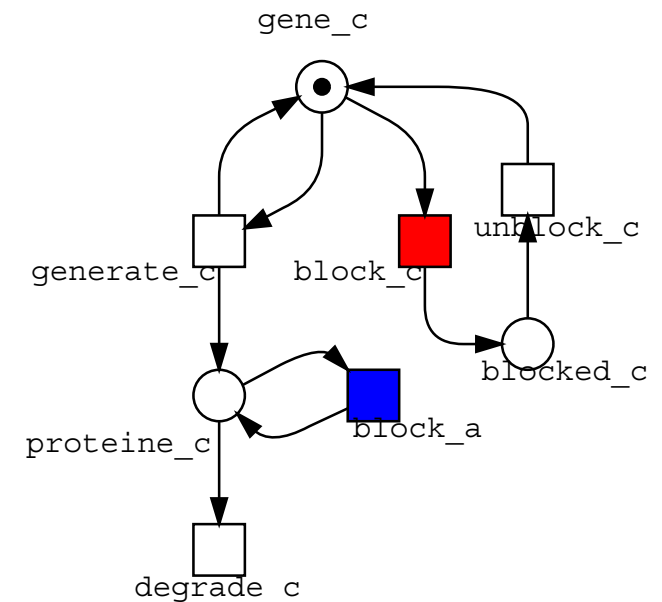
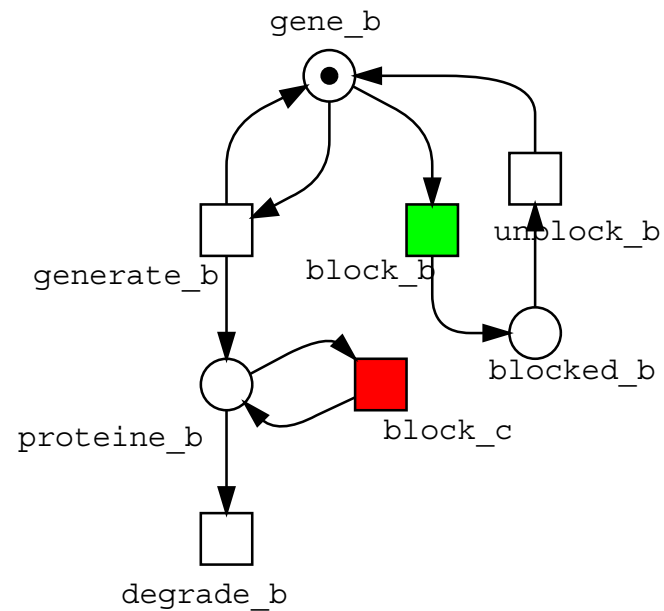
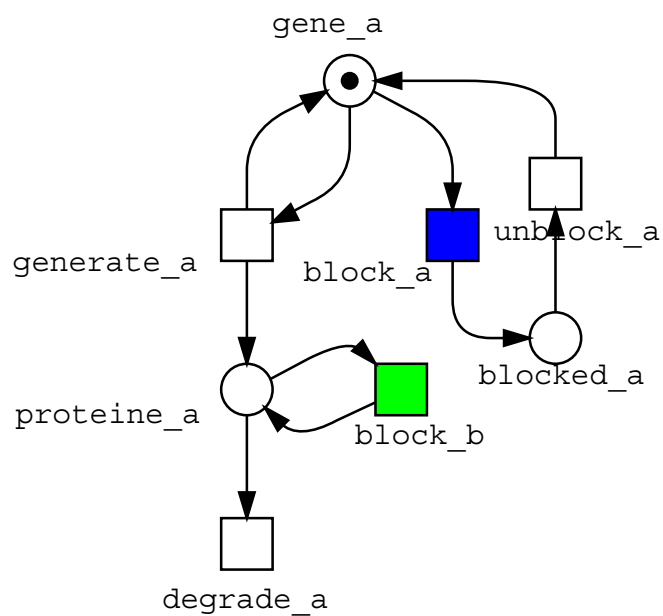


→ Coloured Petri nets – uncoloured Petri nets



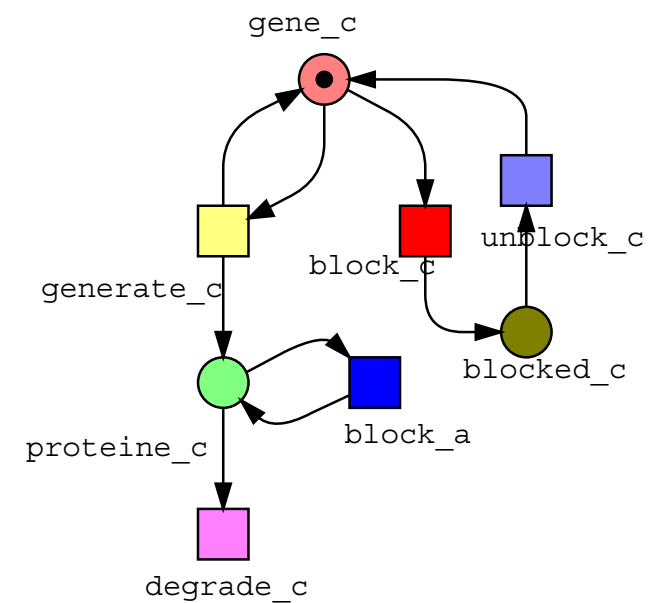
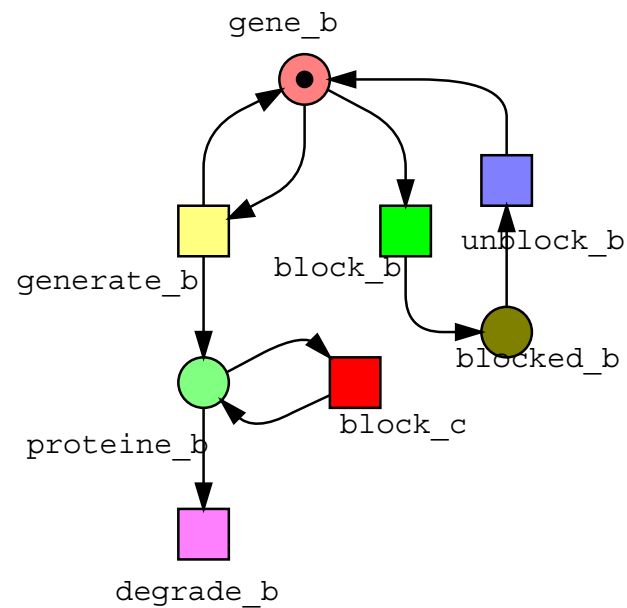
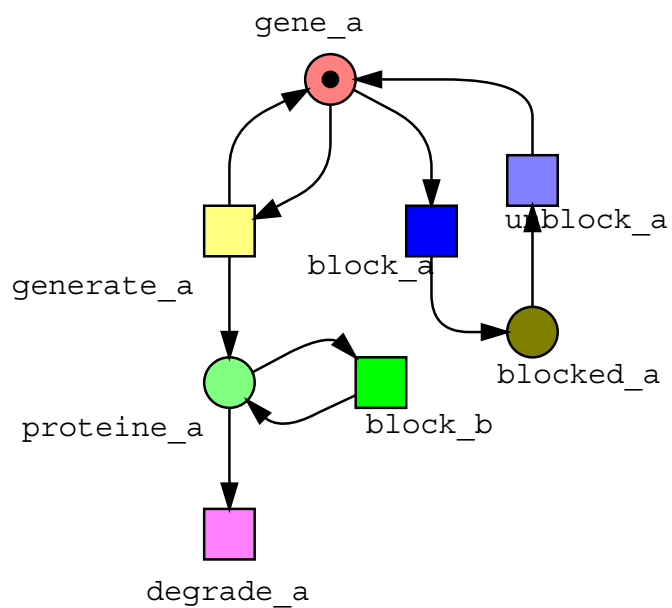
→ Repressilator model

-> Uncoloured Petri nets model



→ Folding repressilator model

- > subnet partition
- > node set partition

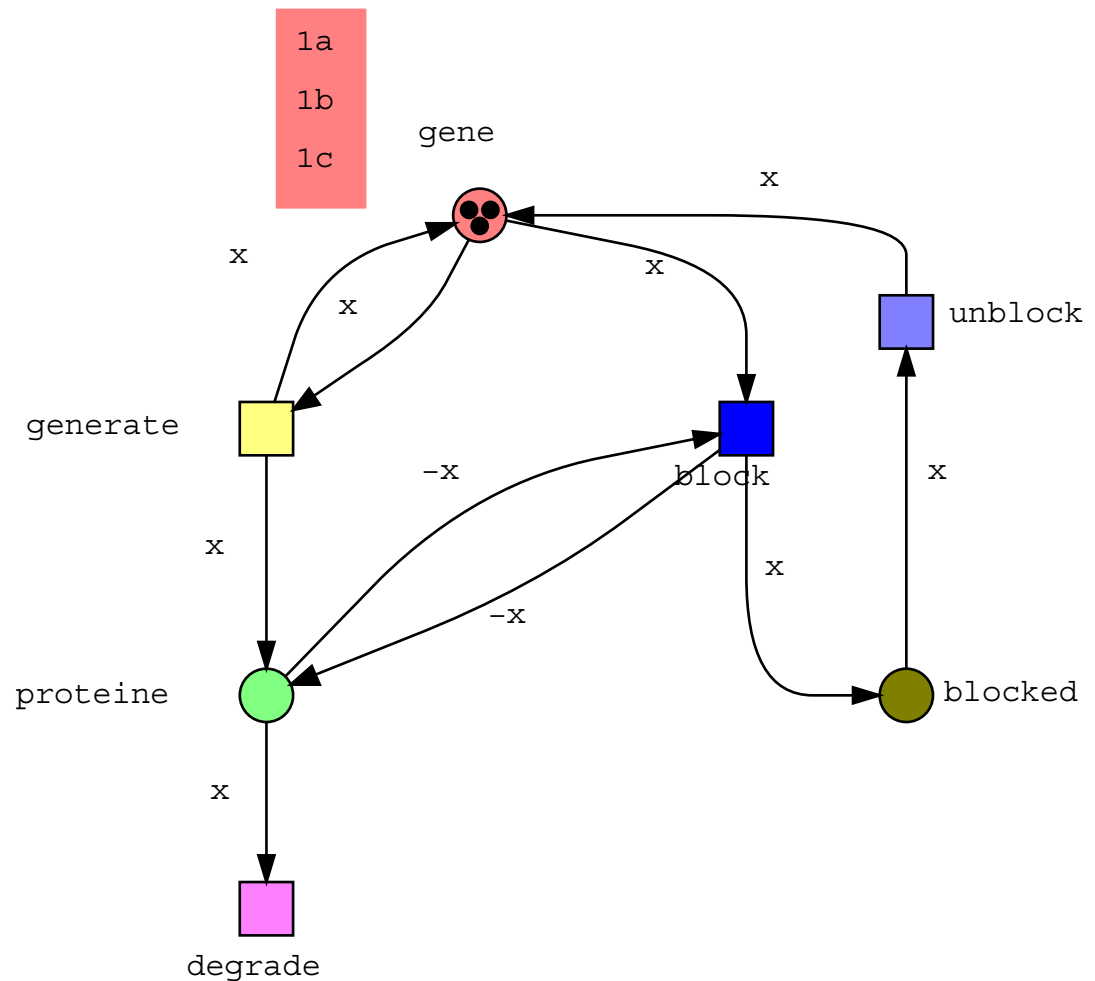


→ Coloured Petri nets for repressilator model

- > define colour sets
- > assign colour sets to places and transitions
- > define arc expressions

Colourset:
enumeration gene = {a, b, c}

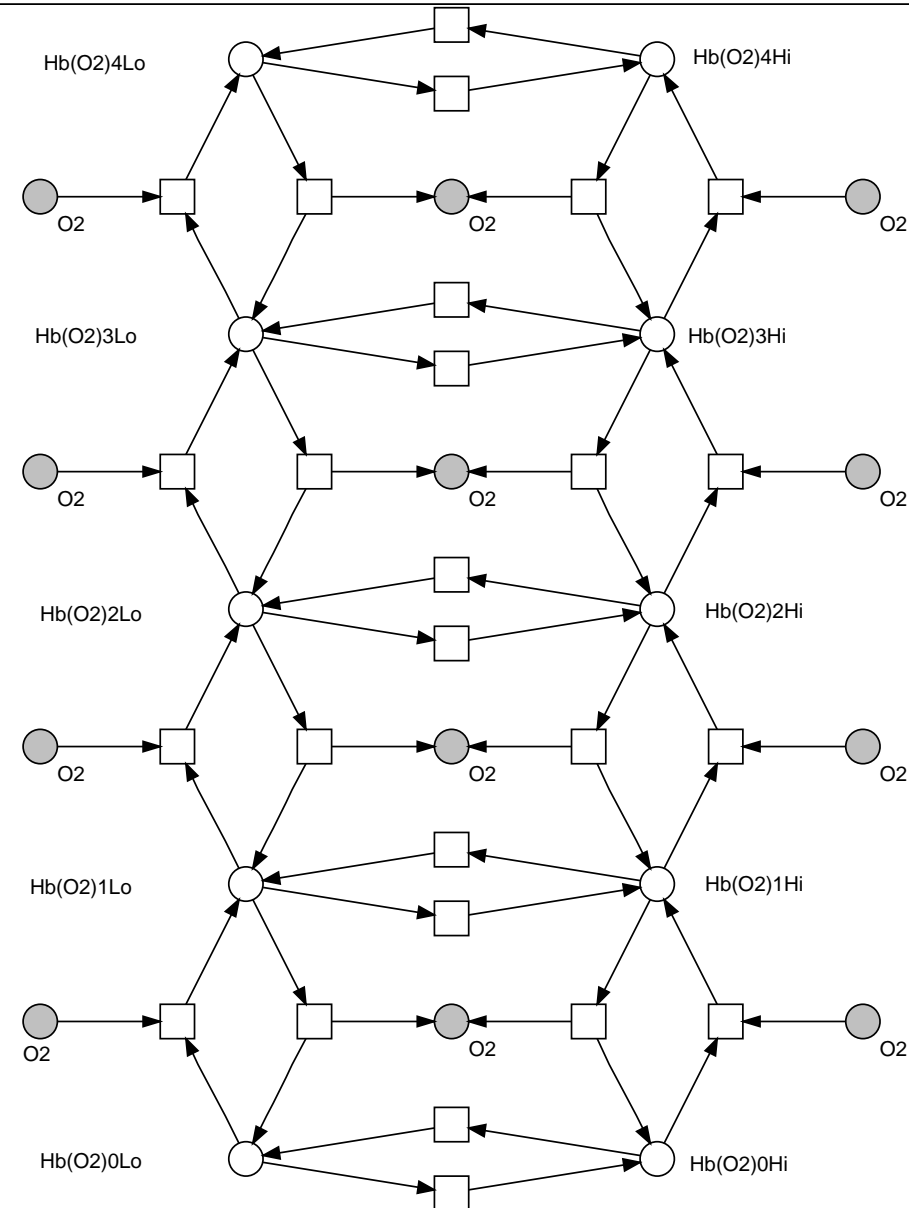
Variable:
gene x



→ Cooperative ligand binding

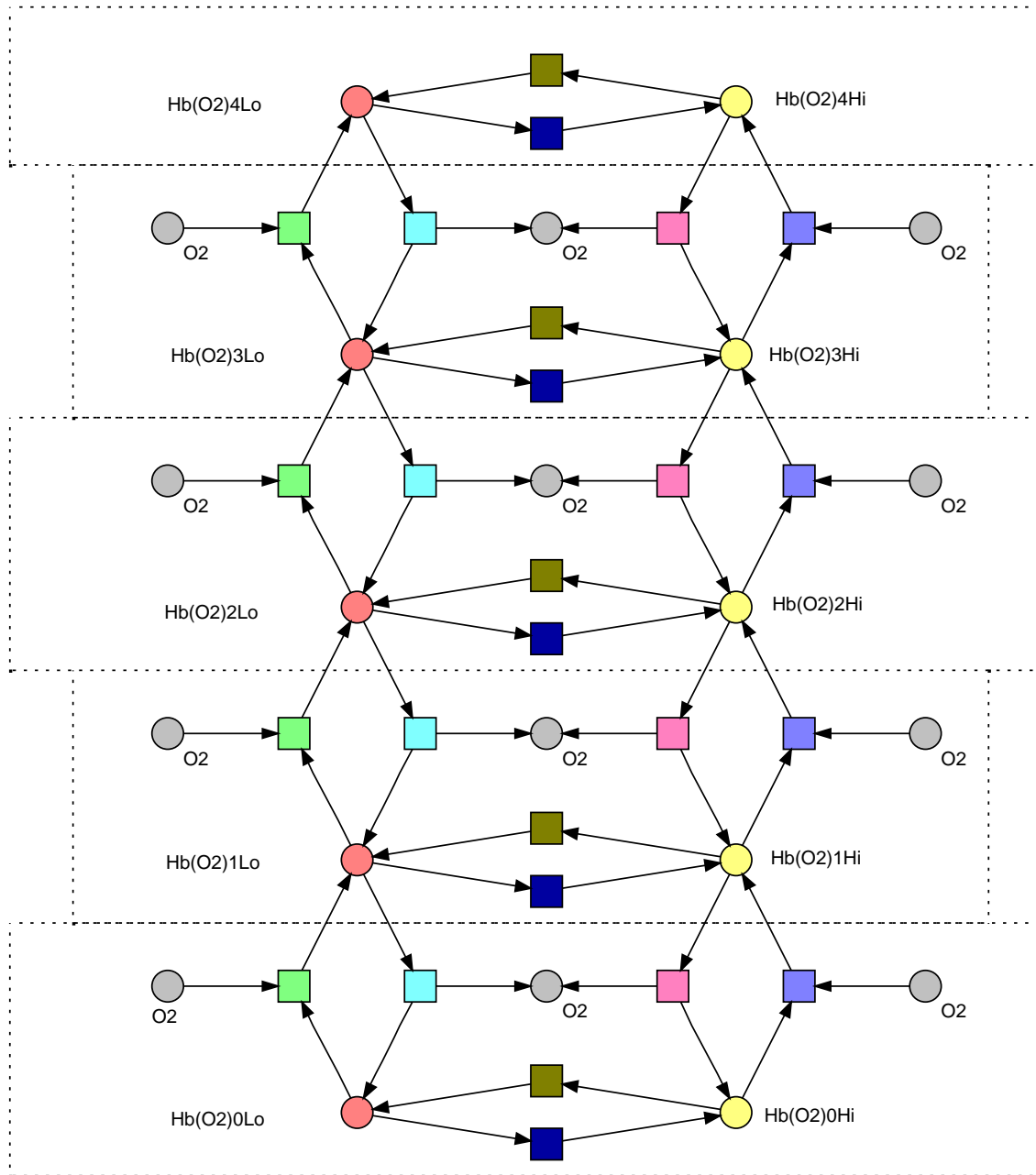
-> Binding of oxygen to the four subunits of a hemoglobin heterotetramer

-> Oxygen (O2) is represented in the form of multiple copies of one logical place



Reference: Annegret Wagler; Robert Weismantel; Wolfgang Marwan. Petri Nets as a Framework for the Reconstruction and Analysis of Signal Transduction Pathways and Regulatory Networks. (so far unpublished)

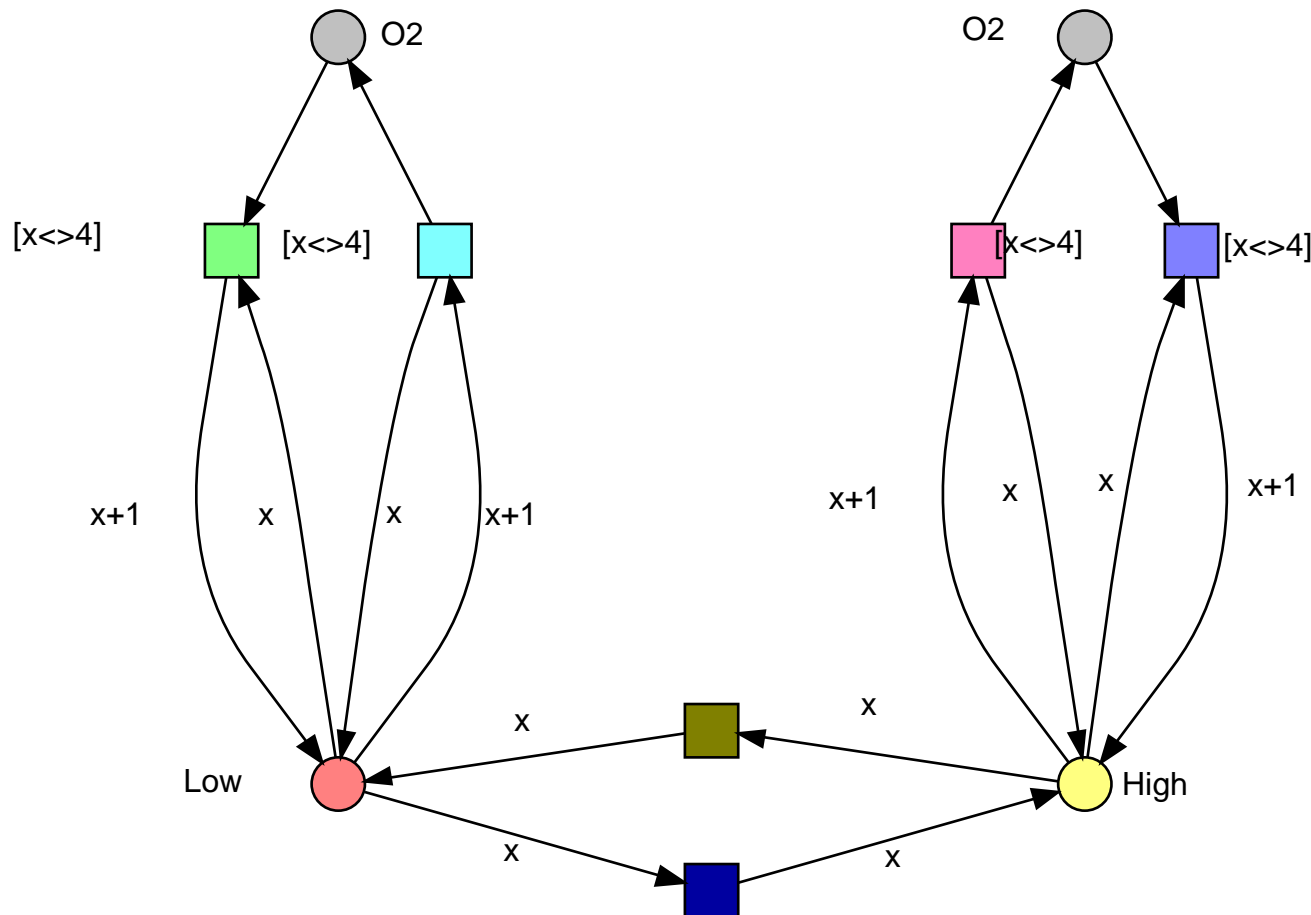
→ Folding



→ Coloured Petri nets

Colourset:
 enumeration $Hb(O_2) = \{Hb(O_2)0, Hb(O_2)1, Hb(O_2)2, Hb(O_2)3, Hb(O_2)4\}$

Variable: $Hb(O_2) \ x$



→ Coloured Petri nets

Colourset:

enumeration $Hb(O2) = \{Hb(O2)0, Hb(O2)1, Hb(O2)2, Hb(O2)3, Hb(O2)4\}$

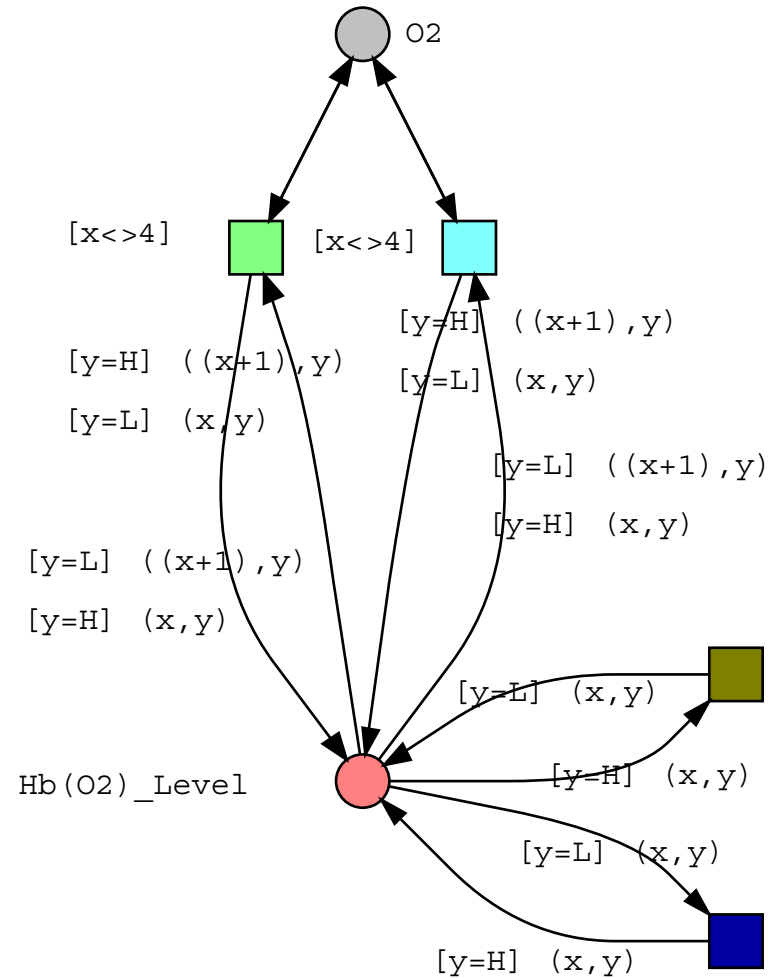
enumeration $Level = \{H, L\}$

Product $Hb(O2)_{Level} = Hb(O2) \times Level$

Variable:

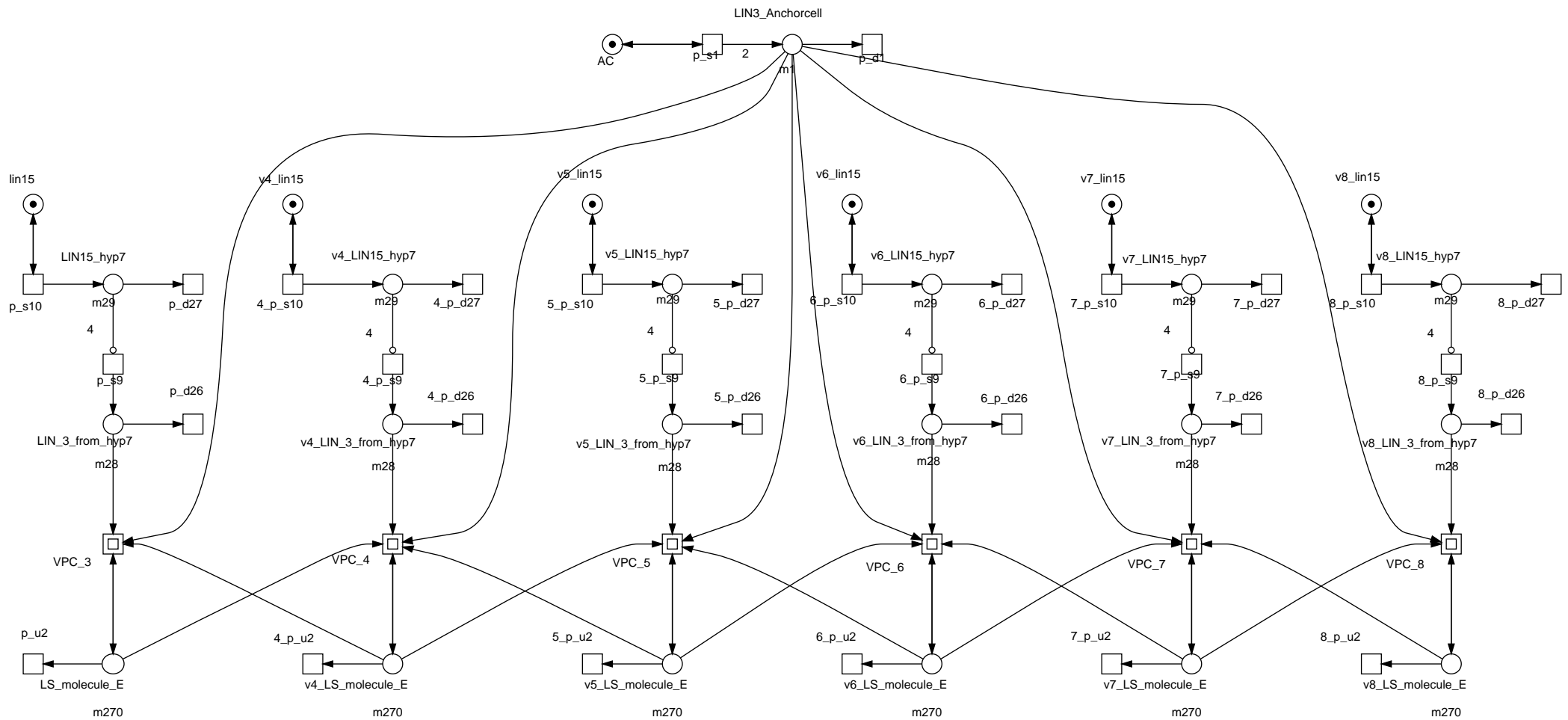
$Hb(O2) \times$

Level y

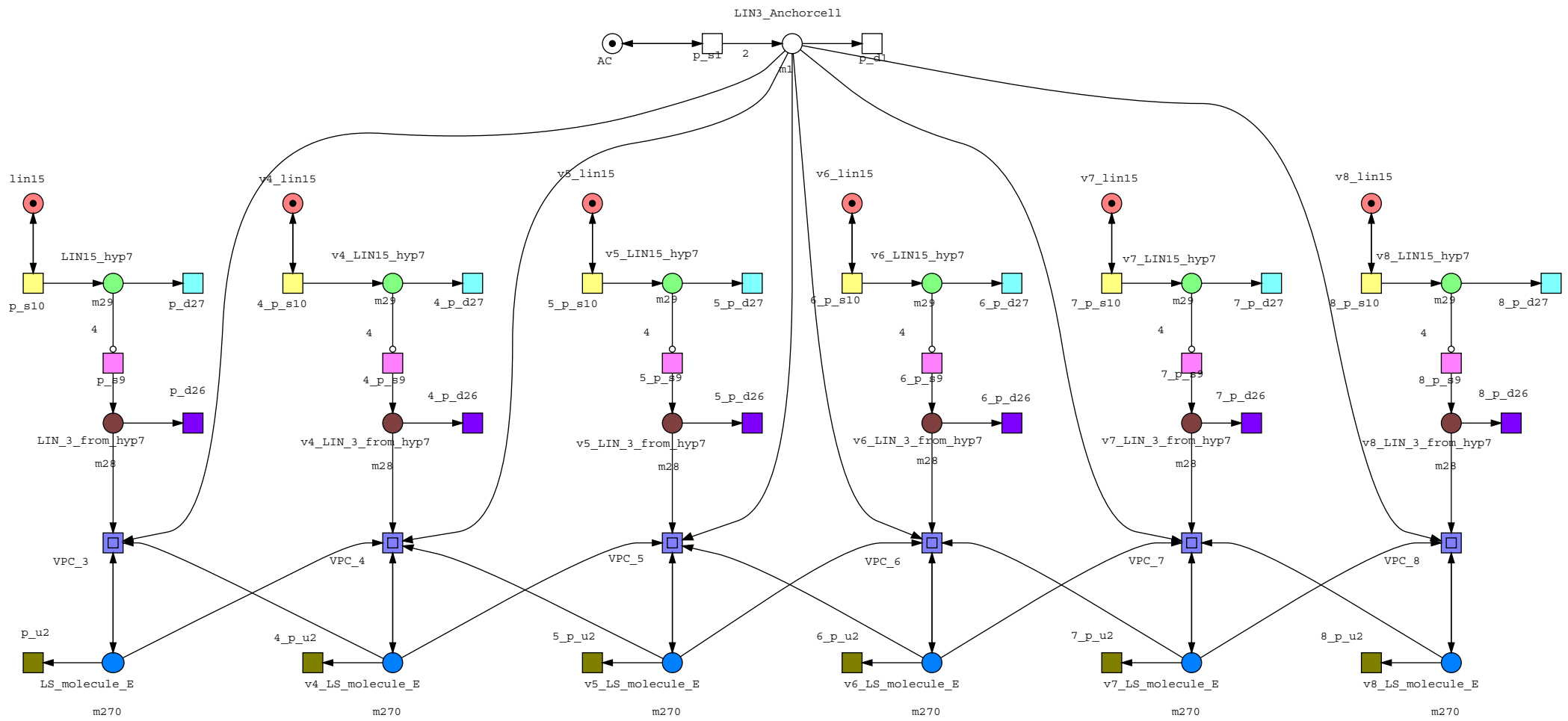


→ C. elegans vulval development model

Places: 206
 Transitions: 360
 Kinetics: mass action



→ Folding



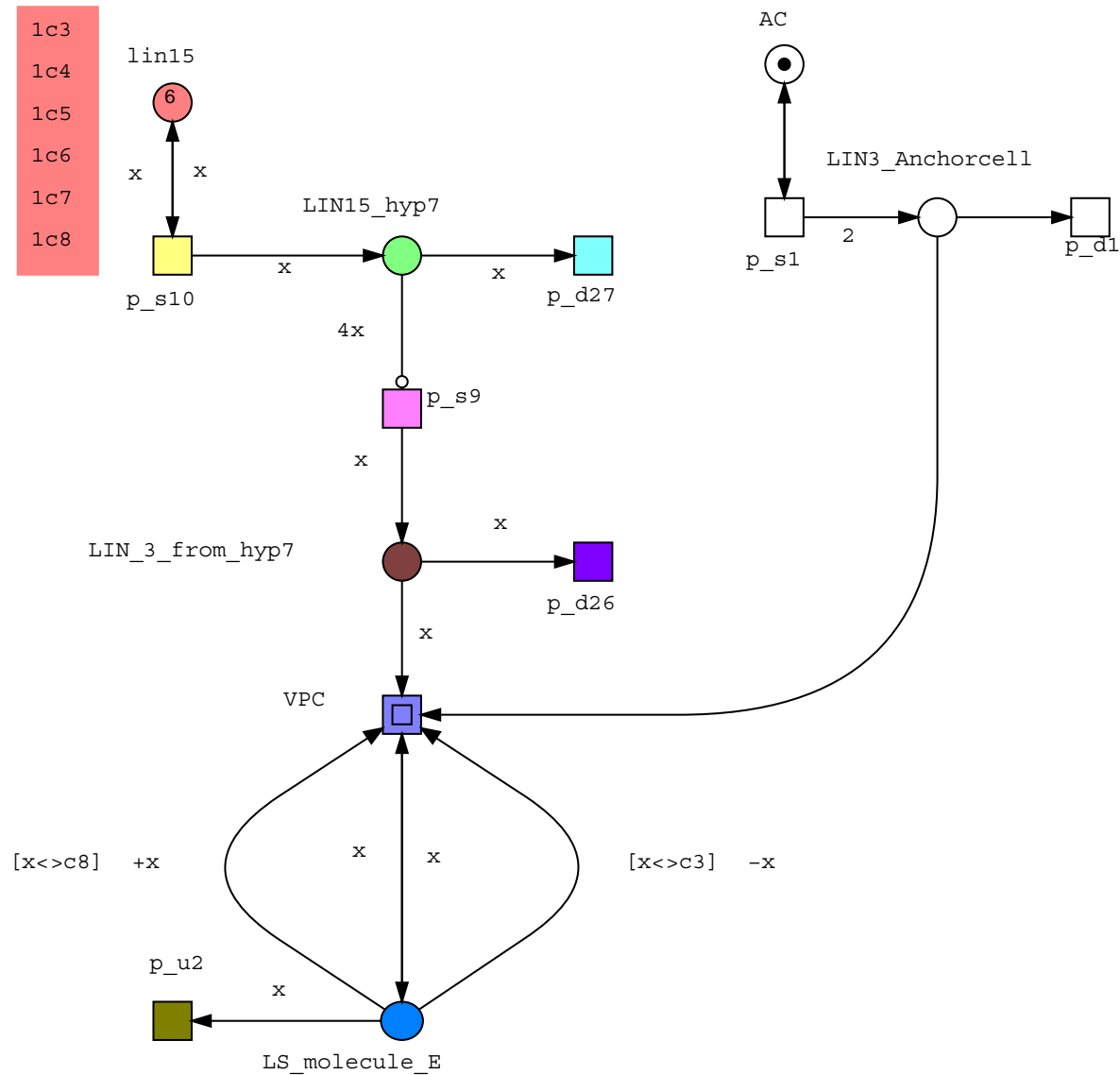
→ C. elegans vulval development model

-> Coloured Petri net

Colourset:
enumeration

Cell={c3, c4, c5, c6, c7, c8}

Variable:
Cell x

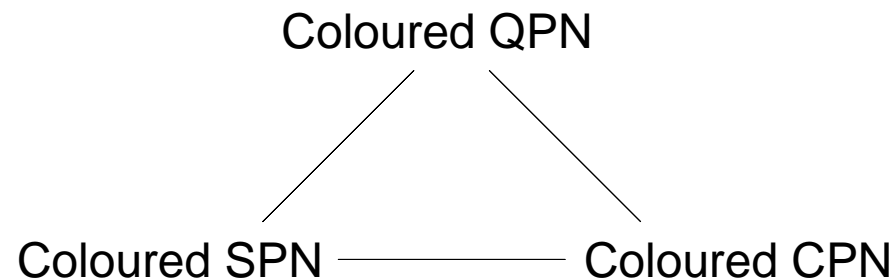


- Reduce the size of nets by folding similar subnets
- To increase the colours means to increase the size of the net
 - > E.g. repressilator model, gene= {a,b,c,...}
- Rich analysis techniques

- by unfolding, reuse all P/T net analysis techniques
 - > invariant analysis
 - > state equation
 - > reachability graph
 - > model checking

- without unfolding, use analysis techniques for coloured Petri nets
 - > reachability graph
 - > model checking

- Requirements analysis for coloured Petri nets
- Precisely defining different types of coloured PN
 - > Coloured qualitative Petri nets (CQPN)
 - > Coloured stochastic Petri nets (CSPN)
 - > Coloured continuous Petri nets (CCPN)
- Defining data types or colours that will be supported
- Defining expressions, guards that will be supported



→ Definitions of data types

<type> ::= <basic type> | <structured type>
<basic type> ::= <type identifier> | <simple type>
<type identifier> ::= <unsigned integer> | <boolean> | <string>
<unsigned integer> ::= "int"
<boolean> ::= "bool"
<string> ::= "string"
<simple type> ::= <enumeration> | <index>
<enumeration> ::= <identifier list>
<identifier list> ::= <identifier> {, <identifier>}
<index> ::= <identifier> "[" <index specifier> "]"
<index specifier> ::= <type identifier> | <enumeration>
<structured type> ::= <product> | <union>
<product> ::= <type> "x" <type> { "x" <type>}
<union> ::= <type> { ", " <type>}

Next Steps

- Develop tools for coloured Petri nets
- Consider more network examples especially from the collaborators
- Continue to analyze the capabilities and equivalence of Cell Illustrator and Snoopy
- Coupling with the database developed by M2, needs to be discussed

Thank You !