P Meta Framework

Assembly of Frequency Dividers

Suprachiasmatic Nucleus

Maintenance of Chronobiological Information by P System Mediated Assembly of Control Units for Oscillatory Waveforms and Frequency

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### Oscillatory Signals in Biology: Range of Periodicities



An individual organism comprises a variety of regulated oscillations.

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### Waveforms of Oscillatory Signals in Biology

Waveform as additional information carrier beyond frequency



Corresponding oscillators for each waveform by small or medium-sized reaction networks

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### 17-years and 13-years Periodical Cicadas with Synchronous Life Cycle



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- No external stimulus with natural period of 17 or 13 years known up to now
- Molecular mechanism to precisely measure the passage of 17 (or 13) years?
- Need of a chemical frequency divider model, ideally configurable for distinct division ratios
- Low number of slight evolutionary changes sufficient to toggle the life cycle between a variety of years?
- ⇒ Assembly of pre-defined chemical modules towards new or extended functionality

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### Inspiration and Motivating Questions

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### 1. Biological Rhythms

### 2. P Meta Framework

- Non-probabilistic P modules
- Connectivity of P modules
- Instructions for composition of P modules
- 3. Assembly of Frequency Dividers
- 4. Suprachiasmatic Nucleus

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### Non-probabilistic P Module

- is able to fulfill an elementary computational task on the fly
- building block of an analog computer or in a control loop
- represents a container encapsulating a formal description of its dynamical behaviour
- specifies the interface of a general real-valued system or its approximation
- aims to bridge building blocks in systems theory and membrane systems

More formally, a P module is a triple  $(\downarrow,\uparrow,\Box)$  where

Each signal is a real-valued function over time.

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 $= (I_1, \dots, I_j) \dots \dots \text{ indicates a list of input signal identifiers}$ =  $(O_1, \dots, O_o) \dots \dots \text{ indicates a list of output signal identifiers}$ ] \dots \d

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### Non-probabilistic P Module as a Processing Unit

input signals



output signals

system providing input-output mapping on the fly

- metabolic P system (mP system) M
- P system for cell signalling modules Π<sub>CSM</sub>
- $\,^{\circ}$  P system for cell signalling networks  $\Pi_{\rm CSN}$
- ordinary differential equations (ODEs) in conjunction with numerical solver
- transfer function (input-output mapping) on its own, given explicitly or implicitly
- characteristic curve, given by numeric values along with approximation/interpolation algorithm

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### P Meta Framework: Connecting P Modules

Our P meta framework is a construct  $\Pi_{\pi\uparrow\downarrow} = (M, P)$  where

M ..... denotes a finite multiset of non-probabilistic P modules P ..... assembly programme by its connectivity instructions



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### P Meta Framework: Connecting P Modules

Our P meta framework is a construct  $\Pi_{\pi\uparrow\downarrow} = (M, P)$  where M ..... denotes a finite multiset of non-probabilistic P modules

P .....assembly programme by its connectivity instructions



- supp(*M*) can be interpreted as the genetic potential of highly conserved reaction units
- Multiplicity of modules → limitation of resources available for module composition

 $V := \{m[i] \mid m \in \operatorname{supp}(M) \land i \in \{1, \dots, M(m)\}\}$  $E := \emptyset$ 

- Indexing of all instances (copies) *m*[*i*] from a P module *m* allows unique identification
- Let a = (a<sub>↓</sub>, a<sub>↑</sub>, a<sub>□</sub>) ∈ supp(M) and b = (b<sub>↓</sub>, b<sub>↑</sub>, b<sub>□</sub>) ∈ supp(M) be two module instances derived from M.
- An edge (a, b, R<sub>a→b</sub>) ∈ E denotes a connection from a to b where dedicated output species of a act as input species of b.
- Each edge comes with a binary relation R<sub>a→b</sub> ⊆ a<sub>↑</sub> × b<sub>↓</sub> in which the mapping of a's output species onto b's input species is given.
- $R_{a \rightarrow b}$  handled in an injective manner since one output species is allowed to cover several downstream input species, but each input species must be supplied by at most one upstream output species.

### P System Mediated Assembly of Control Units in Chronobiology

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Assembly of Frequency Dividers

Suprachiasmatic Nucleus

### Assembly Programme Instructions (Selection)

- Time-dependent changes of module connectivity
- Time stamp *t* opens each instruction
- Instruction updates graph G(V, E)
- t: ModuleConnect $(a \rightarrow b, R_{a \rightarrow b})$

connects some or all of module *a*'s output species to represent *b*'s input species by sharing species identifiers according to the injective binary relation  $R_{a\to b} \subseteq a_{\uparrow} \times b_{\downarrow}$ .

Edge update scheme:  $E := E \cup \{(a, b, R_{a \rightarrow b})\}$ 



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### Assembly Programme Instructions (Selection)

- Time-dependent changes of module connectivity
- Time stamp *t* opens each instruction
- Instruction updates graph G(V, E)
- t : ModuleDisconnect $(a \leftrightarrow b)$

completely disconnects modules *a* and *b* by annihilating all cross-modular species sharings. This comes along with removing  $R_{a\rightarrow b}$  as well as  $R_{b\rightarrow a}$ , respectively.

Edge update scheme:  $E := E \setminus \{(a, b, R_{a \rightarrow b})\} \setminus \{(b, a, R_{b \rightarrow a})\}$ 

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- 1. Biological Rhythms
- 2. P Meta Framework
- 3. Assembly of Frequency Dividers
  - Chemical frequency divider model 1:17
  - Frequency divider 1:5 by separator removal
  - Frequency divider 1:6 by Repressilator
  - Frequency divider 1:3 by Goodwin module
- 4. Suprachiasmatic Nucleus

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## The Repressilator Module

### repressilator = $(\emptyset, \{Z\}, F)$



- Oscillation by progressional inhibition in gene regulatory network
- Involves higher-order Hill kinetics
- Velocity of decay reactions mainly determines frequency
- Inherently almost sinusoidal waveform

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## The Goodwin Module

### goodwin = $(\emptyset, \{X\}, F)$



- Oscillation by mutual activation and inhibition in gene regulatory network
- Involves higher-order Hill kinetics
- Velocity of decay reactions mainly determines frequency
- Configurable to exhibit almost sinusoidal or plated waveforms

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# The Brusselator Module brusselator = $(\emptyset, \{S\}, F)$



- Oscillation by autocatalytic feedback loops
- Exclusively mass-action kinetics
- Velocity of decay in concert with autocatalytic loop determines frequency
- Inherently spiking waveform

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# The Binary Signal Separator Module separator = $({O_0^F}, {O_3^F}, F)$



- · Converts spiking or sinusoidal oscillations into plated waveform
- Successive binarisation by enzymatically controlled cascade
- Exclusively mass-action kinetics

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### Binary Counter Modulo 17 – Chemical Logic gate

Boolean variable *z* represented by two correlated species  $Z^T$  and  $Z^F$ 

### **Chemical reaction network for NAND**



### Analogously implementation of other boolean functions

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### The Binary Counter Modulo 17 $mod17 = (\{C\}, \{B_1^T\}, F)$



- Based on 5-bit Gray code
- Clock signal C increments counter
- Exclusively mass-action kinetics

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### Composing the Original Frequency Divider 1:17



 $\begin{aligned} \Pi_{\text{FD17}} &= (M, P) \quad \text{with} \\ M &= \{(\text{brusselator}, 1), (\text{repressilator}, 1), (\text{goodwin}, 1), (\text{separator}, 1), (\text{mod}17, 1)\} \\ P &= \{0 : \text{ModuleConnect}(\text{brusselator}[1] \rightarrow \text{separator}[1], \{(S, O_0^F)\}), \\ 0 : \text{ModuleConnect}(\text{separator}[1] \rightarrow \text{mod}17[1], \{(O_3^F, C)\})\} \end{aligned}$ 

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### Frequency Divider 1:5 by Removal of Signal Separator



$$\begin{split} \Pi_{\text{FD5}} &= (M, P) \quad \text{with} \\ M &= \{(\text{brusselator}, 1), (\text{repressilator}, 1), (\text{goodwin}, 1), (\text{separator}, 1), (\text{mod}17, 1)\} \\ P &= \{0 : \text{ModuleConnect}(\text{brusselator}[1] \rightarrow \text{separator}[1], \{(S, O_0^F)\}), \\ 0 : \text{ModuleConnect}(\text{separator}[1] \rightarrow \text{mod}17[1], \{(O_3^F, C)\}), \\ 200 : \text{ModuleDisconnect}(\text{brusselator}[1] \rightarrow \text{mod}17[1], \{(S, C)\})\} \\ 200 : \text{ModuleConnect}(\text{brusselator}[1] \rightarrow \text{mod}17[1], \{(S, C)\})\} \end{split}$$

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### Frequency Divider 1:6 by Repressilator



 $\begin{aligned} \Pi_{\text{FD6}} &= (M, P) \quad \text{with} \\ M &= \{(\text{brusselator}, 1), (\text{repressilator}, 1), (\text{goodwin}, 1), (\text{separator}, 1), (\text{mod}17, 1)\} \\ P &= \{0 : \text{ModuleConnect}(\text{repressilator}[1] \rightarrow \text{mod}17[1], \{(Z, C)\})\} \end{aligned}$ 

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### Frequency Divider 1:3 by Goodwin Module



 $\begin{aligned} \Pi_{\text{FD3}} &= (M, P) \quad \text{with} \\ M &= \{(\text{brusselator}, 1), (\text{repressilator}, 1), (\text{goodwin}, 1), (\text{separator}, 1), (\text{mod}17, 1)\} \\ P &= \{0 : \text{ModuleConnect}(\text{goodwin}[1] \rightarrow \text{mod}17[1], \{(X, C)\})\} \end{aligned}$ 

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- 1. Biological Rhythms
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- 4. Suprachiasmatic Nucleus
  - Core Oscillator's Interplay in Suprachiasmatic Nucleus
  - Prospectives
  - Acknowledgement

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### Core Oscillator's Interplay in Suprachiasmatic Nucleus



 $\Pi_{\rm SCN} = (M, P)$  with

 $M = \{(n, 14)\}$ 

 $P = \{0 : ModuleConnect(n[1] \rightarrow n[3], \{(Y, X)\}),\$ 

- $\texttt{0} : \texttt{ModuleConnect}(n[1] \rightarrow n[4], \{(\textbf{\textit{Y}}, \textbf{\textit{X}})\}), \ldots,$
- $\texttt{0} : \texttt{ModuleConnect}(\texttt{n[10]} \rightarrow \texttt{n[14]}, \{(\texttt{Y}, \texttt{X})\})$
- $\texttt{300} : \texttt{ModuleConnect}(n[2] \rightarrow n[4], \{(\textbf{\textit{Y}}, \textbf{\textit{X}})\}), \ldots,$
- $\texttt{300} : \texttt{ModuleConnect}(\texttt{n[6]} \rightarrow \texttt{n[10]}, \{(Y, X)\})\}$

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### Prospectives

### Conclusions

- Assembly and reassembly as well as composition and decomposition of pre-defined reaction network modules on the fly —> promising strategy in order to achieve complex systems capable of new, extended, or unexpected functionality
- Framework for an artificial evolution at a granularity of highly conserved genetic ensembles
- P system-based approach features by coping with dynamical structures at a modular level rather than a molecular level
- Simulation studies carried out using CoPaSi along with Gepasi Model Extractor

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### **Further work**

- Incorporation of Infobiotics workbench
- Extension of P module library
- Combining advantages of P system-based approaches with analytical examination tools using ODEs

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### Conclusions

- Assembly and reassembly as well as composition and decomposition of pre-defined reaction network modules on the fly —> promising strategy in order to achieve complex systems capable of new, extended, or unexpected functionality
- Framework for an artificial evolution at a granularity of highly conserved genetic ensembles
- P system-based approach features by coping with dynamical structures at a modular level rather than a molecular level
- Simulation studies carried out using CoPaSi along with Gepasi Model Extractor

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- Combining advantages of P system-based approaches with analytical examination tools using ODEs

P System Mediated Assembly of Control Units in Chronobiology

P Meta Framework

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Suprachiasmatic Nucleus

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### Acknowledgements





Jena Centre for Bioinformatics

## b-tu

Brandenburg University of Technology Cottbus

DEUTSCHE FORSCHUNGSgemeinschaft, grant Hi801/3-1

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