

Reversibility in massive concurrent systems

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MecBic 2011

programme

- study reversibility in DNA circuits
- define the algebra of DNA circuits – *reversible structures*
- analyze the interplay between reversibility and causal dependency
 - + grasp causal dependency between coinital reductions
 - + investigate causal equivalent computations (permutation equivalence)
- measure the expressive power of reversible structures
 - + implementation of asynchronous RCCS

“natural” reversibility?

- in computational systems, computations are sequence of *irreversible steps*
- implementations of these systems in physics or chemistry are usually reversible
- reversibility means undoing the computation *not in a deterministic way*:

states reached during a backward computation are states that could have been reached during the forward computation by just performing independent actions in a different order

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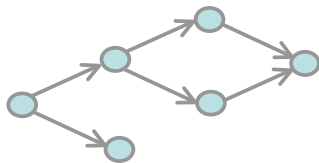
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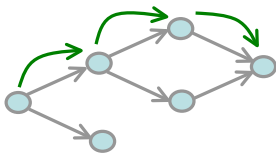
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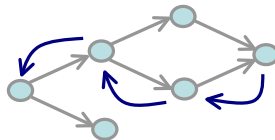
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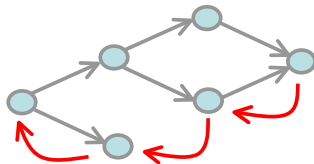
transition system



a forward computation

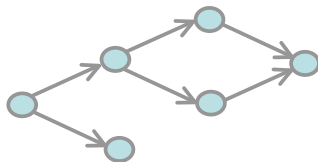


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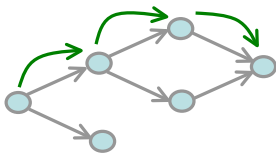


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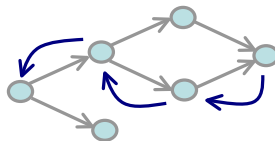
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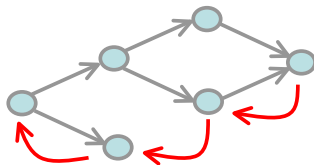
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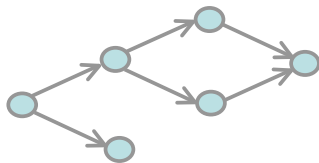


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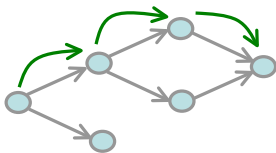


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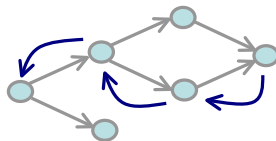
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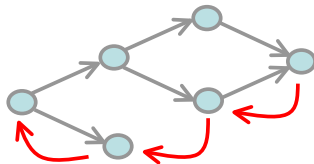
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since part of physics and chemistry is reversible,

what is the formal theory of reversibility in these fields?

said otherwise:

taking a natural reversible system, what properties may we prove?

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reversibility in DNA circuits (cf. Cardelli/Phillips)

- subsequences on a DNA strand are called **domains**



domains are **independent** of each other

- they cannot hybridize from any other domain except their complement
- there are very few short domains with reversible hybridizations
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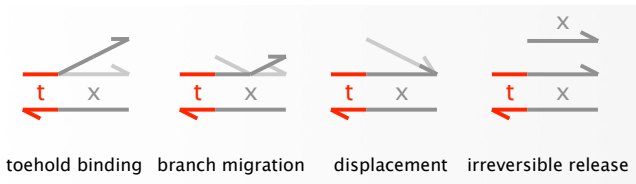
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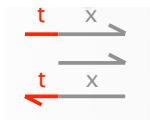
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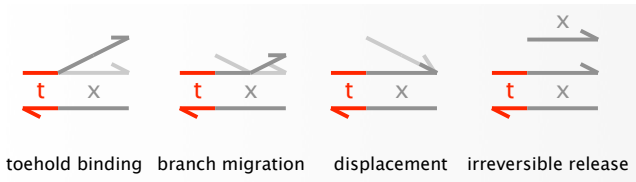
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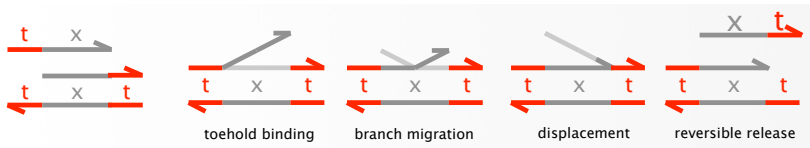
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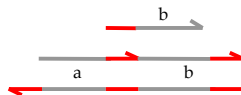


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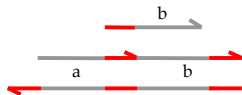
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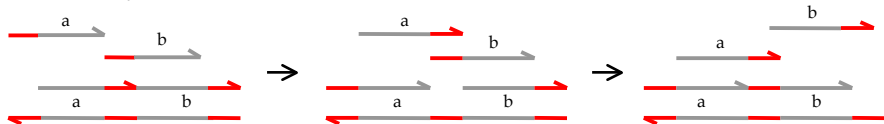
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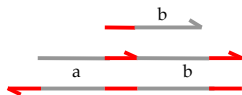
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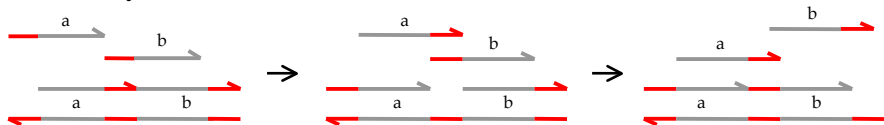
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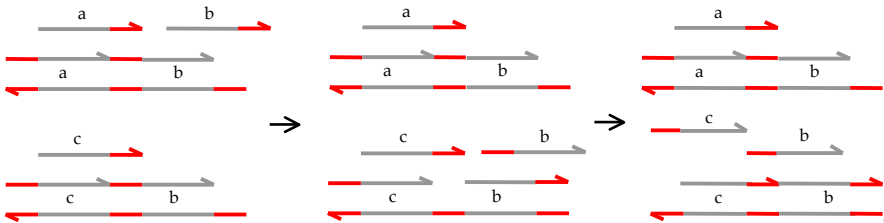
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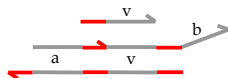


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a better model for causality: three-domains structures

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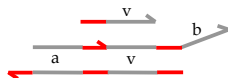


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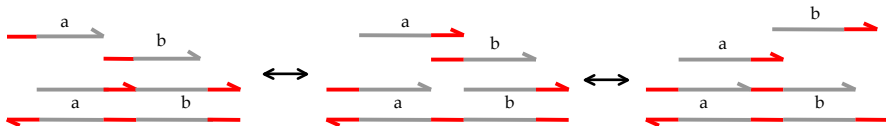
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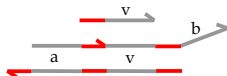
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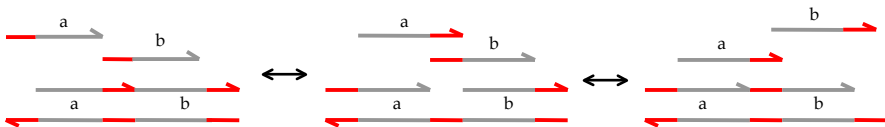
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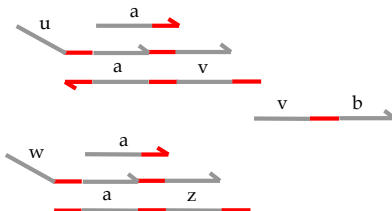
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causality and massive concurrency

DNA circuits are *massively concurrent*:

- solutions consist of populations of species of strands and
- populations are not singletons

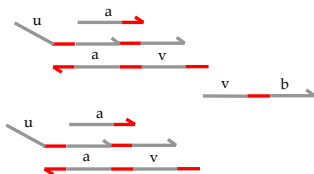
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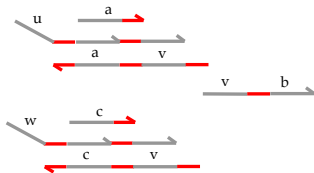
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the situation may be even worse due to bad designs



a theory of massive concurrency?

reversibility/causality in massive concurrent systems has not been studied

- theories have been defined for reversible calculi where processes retain unique ids (Danos-Krivine, Phillips-Ulidowski, Lanese-Mezzina-Stefani)

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the algebra of reversible structures

notation: u, v, w : ids
 a, \bar{a}, b, \bar{b} : names and conames
 x, x' : ids, names and conames
 A, B, C : sequences of names;
 $\bar{A}, \bar{B}, \bar{C}$: sequences of elements $u : \bar{a}$
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signals : $u : \bar{a}$

gates : g input part.output part + $\hat{}$

examples of gates:

$\hat{a} . a' . v : \bar{b}$ $u : a . \hat{a}' . v : \bar{b}$

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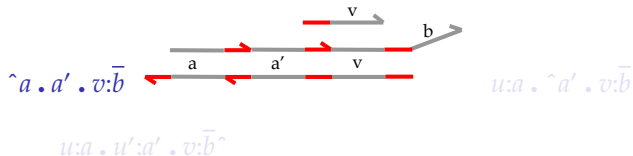
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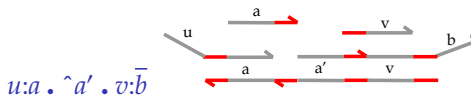
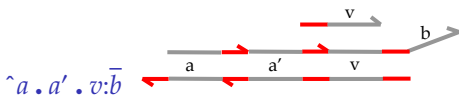
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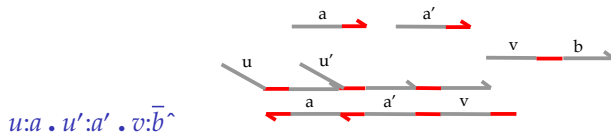
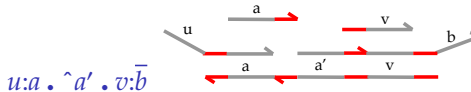
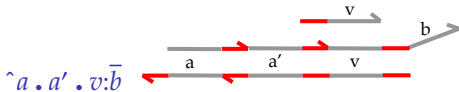
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the algebra of reversible structures – syntax

structures : $S ::=$

	$\mathbf{0}$	(null)
	$u:\bar{a}$	(signal)
	g	(gate)
	$S \mid S$	(parallel)
	$(\text{new } x)S$	(new)

the algebra of reversible structures – reductions

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$$v:\bar{a} \mid w:\bar{a} \mid \hat{a} . u:\bar{b} \begin{array}{l} \nearrow w:\bar{a} \mid v:a . \hat{u}:\bar{b} \rightarrow w:\bar{a} \mid v:a . u:\bar{b} \hat{} \mid u:\bar{b} \\ \searrow v:\bar{a} \mid w:a . \hat{u}:\bar{b} \end{array}$$

the algebra of reversible structures – reductions

$$\text{input capture: } u:\bar{a} \mid A^\perp . \hat{a} . B . \bar{C} \longrightarrow A^\perp . u:a . \hat{B} . \bar{C}$$

$$\text{input release: } A^\perp . u:a . \hat{B} . \bar{C} \longrightarrow u:\bar{a} \mid A^\perp . \hat{a} . B . \bar{C}$$

$$\text{output release: } A^\perp . \bar{B} . \hat{u}:\bar{a} . \bar{C} \longrightarrow u:\bar{a} \mid A^\perp . \bar{B} . u:\bar{a} . \hat{C}$$

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pearls of expressive power

join input ($a \mid b \triangleright \bar{c}$)

$\hat{a}.b.u:\bar{c}$

input-guarded choice ($a.\bar{b} + a'.\bar{c}$)

$(\text{new } v, e)(e.a.u:\bar{b} \mid e.a'.u':\bar{c} \mid v:\bar{e})$

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weak coherence

remark: it is possible to restrict the arguments about the dynamics of reversible structures to structures without news (*this simplifies the theory*)

a structure S is weak coherent whenever ids are uniquely associated to names and co-names

(if $u : \alpha$ and $u : \alpha'$ occur in S' then either $\alpha = \alpha'$ or $\alpha = \overline{\alpha'}$)

proposition: *weak coherent reversible algebra may be implemented into three domains DNA circuits*

– the correspondence is consistent and complete wrt reductions

causality: the tour

1. define *causal dependence* on coinital reductions
 - two reductions are dependent if they have either the signal or the gate in common
 - independent reductions can be swapped (diamond lemma)
2. define *permutation equivalence*, an equivalence on derivations that is insensible
 - i. to swapping of independent reductions
 - ii. to the removal of reverse reductions
3. study the theory of permutation equivalence

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causal independence

a reduction can be addressed by the signal/gate that move

remark: *in weak-coherent structures, names and co-names when prefixed by ids are useless*

– a signal $u : \bar{a}$ can be addressed by u

– a gate

$\hat{a}.a'.v : \bar{b}$ can be addressed by $\hat{a}a' \circ v$

$u : a.\hat{a}'.v : \bar{b}$ can be addressed by $u\hat{a}' \circ v$

$u : a.u' : a'.v : \bar{b}$ can be addressed by $uu' \circ v\hat{a}$

cf. Lévy labels in lambda calculus

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causal independence/cont.

an example

$$\begin{array}{ccc} u:\bar{a} \mid \hat{a} \cdot v:\bar{b} \mid u:a \cdot \hat{v}:\bar{b} & & \\ \color{red} u \mid \hat{a} \circ v \swarrow & & \searrow u \circ \hat{v} \\ u:a \cdot \hat{v}:\bar{b} \mid u:a \cdot \hat{v}:\bar{b} & & u:\bar{a} \mid \hat{a} \cdot v:\bar{b} \mid u:a \cdot v:\bar{b} \hat{} \mid v:\bar{b} \\ \color{red} u \circ \hat{v} \searrow & & \swarrow u \mid \hat{a} \circ v \\ u:a \cdot \hat{v}:\bar{b} \mid u:a \cdot v:\bar{b} \hat{} \mid v:\bar{b} & & \end{array}$$

two coinital reductions are causally independent if the corresponding labels have no sublabel in common
(causal dependency is the opposite notion)

causal independence/cont.

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the diamond lemma

Lemma. *Let $S \xrightarrow{\mu} S'$ and $S \xrightarrow{\nu} S''$ be such that μ and ν are causally independent. Then there exists S''' such that $S' \xrightarrow{\nu} S'''$ and $S'' \xrightarrow{\mu} S'''$.*

causal independence: issues

$$\begin{array}{ccc} u:\bar{a} \mid u:\bar{a} \mid \hat{a} \cdot v:\bar{b} \mid w:c \cdot u:\bar{a}^{\wedge} & \xrightarrow{u \mid \hat{a} \circ v} & u:\bar{a} \mid u:a \cdot \hat{v}:\bar{b} \mid w:c \cdot u:\bar{a}^{\wedge} \\ & \xrightarrow{u \mid w \circ u^{\wedge}} & u:a \cdot \hat{v}:\bar{b} \mid w:c \cdot \hat{u}:\bar{a} \end{array}$$

cannot be swapped because $u \mid \hat{a} \circ v$ and $u \mid w \circ u^{\wedge}$ have sublabels in common

rationale:

- labels are not expressive enough to catch *multiplicities*
 - a similar anomaly is present in Petri nets (cf. Degano, Meseguer, Montanari)
- *in massive concurrent systems, different occurrences of a same molecule cannot be separated*

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permutation equivalence

notation:

let $\mu_1; \dots; \mu_n$ be the computation $S_1 \xrightarrow{\mu_1} \dots \xrightarrow{\mu_n} S_{n+1}$

let $[\mu]^+$ be the *reverse label* of μ defined as

$$\begin{aligned} [u | v \hat{a} \circ w]^+ &= \tilde{v} u \hat{\circ} \tilde{w} \\ [v u \hat{\circ} w]^+ &= u | v \hat{a} \circ w \\ [v u \circ \hat{w}]^+ &= u | v u \circ w \hat{} \\ [u | v u \circ w \hat{a}]^+ &= v u \circ \hat{w} \end{aligned}$$

permutation equivalence \sim is the least equivalence relation between computations closed under composition and such that:

$$\begin{aligned} \mu; [\mu]^+ &\sim \varepsilon \\ \mu; \nu &\sim \nu; \mu \quad \text{if } \mu \text{ and } \nu \text{ are coinital and causally independent} \end{aligned}$$

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permutation equivalence: examples

$$- \quad v:\bar{a} \mid w:\bar{a} \mid \hat{a}.u:\bar{b} \xrightarrow{v|\hat{a}\circ u} w:\bar{a} \mid v:a.\hat{u}:\bar{b} \xrightarrow{v\hat{\circ}u} v:\bar{a} \mid w:\bar{a} \mid \hat{a}.u:\bar{b}$$

is computationally equivalent to ε

$$- \quad v:\bar{a} \mid w:\bar{a} \mid \hat{a}.u:\bar{b} \mid \hat{a}.z:\bar{c} \xrightarrow{v|\hat{a}\circ u} w:\bar{a} \mid v:a.\hat{u}:\bar{b} \mid \hat{a}.z:\bar{c}$$

$$\xrightarrow{w|\hat{a}\circ z} v:a.\hat{u}:\bar{b} \mid w:a.\hat{z}:\bar{c}$$

$$\xrightarrow{v\hat{\circ}u} v:\bar{a} \mid \hat{a}.u:\bar{b} \mid w:a.\hat{z}:\bar{c}$$

is equivalent to

$$v:\bar{a} \mid w:\bar{a} \mid \hat{a}.u:\bar{b} \mid \hat{a}.z:\bar{c} \xrightarrow{w|\hat{a}\circ z} v:\bar{a} \mid \hat{a}.u:\bar{b} \mid w:a.\hat{z}:\bar{c}$$

$$\xrightarrow{v|\hat{a}\circ u} v:a.\hat{u}:\bar{b} \mid w:a.\hat{z}:\bar{c}$$

$$\xrightarrow{v\hat{\circ}u} v:\bar{a} \mid \hat{a}.u:\bar{b} \mid w:a.\hat{z}:\bar{c}$$

and to

$$v:\bar{a} \mid w:\bar{a} \mid \hat{a}.u:\bar{b} \mid \hat{a}.z:\bar{c} \xrightarrow{w|\hat{a}\circ z} v:\bar{a} \mid \hat{a}.u:\bar{b} \mid w:a.\hat{z}:\bar{c}$$

permutation equivalence: examples

$$- \quad v:\bar{a} \mid w:\bar{a} \mid \hat{a} . u:\bar{b} \xrightarrow{v \mid \hat{a} \circ u} w:\bar{a} \mid v:a . \hat{u}:\bar{b} \xrightarrow{v \mid \hat{a} \circ u} v:\bar{a} \mid w:\bar{a} \mid \hat{a} . u:\bar{b}$$

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$$\xrightarrow{w \mid \hat{a} \circ z} v:a . \hat{u}:\bar{b} \mid w:a . \hat{z}:\bar{c}$$

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permutation equivalence: examples

$$- \quad v:\bar{a} \mid w:\bar{a} \mid \hat{a}.u:\bar{b} \xrightarrow{v|\hat{a}\circ u} w:\bar{a} \mid v:a.\hat{u}:\bar{b} \xrightarrow{v\hat{\circ}u} v:\bar{a} \mid w:\bar{a} \mid \hat{a}.u:\bar{b}$$

is computationally equivalent to ε

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the standardization theorem

let $\mu_1 ; \dots ; \mu_n$ be a computation of a weak coherent structure
such that μ_n is the converse of μ_1

- there is a shorter computation that is permutation equivalent to $\mu_1 ; \dots ; \mu_n$

the evolution of a gate in a computation without converse labels (normal)
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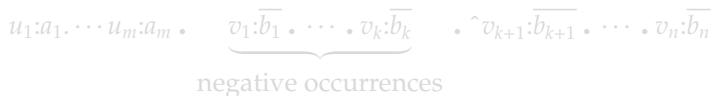
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coherence

a solution must contain exactly one molecule of every species

a weak-coherent structure is *coherent* whenever

- different gates have types with no id in common – *the type of a gate is the sequence of ids in the output part*
- ids occur at most twice: one occurrence is positive and the other is negative



examples: $u:\bar{a} \mid \hat{a} \cdot v:\bar{a}$ and $v:\bar{a} \mid u:a \cdot v:\bar{a}$ are coherent

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consequences of coherence

theorem: *two coinital computations of a coherent structure are permutation equivalent if and only if they are cofinal*

(false in weak-coherent structures)

theorem: *the reachability problem in coherent structure has a computational complexity of $O(n^2)$, where n is the number of gates in the structure*

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expressive power of coherent structures

coherent structures encode in a causally consistent way
asynchronous Reversible CCS

asynchronous Reversible CCS in a nutshell

memories $m ::= \langle \rangle \mid \langle i \rangle_n \bullet m \mid \langle m, \alpha, Q \rangle \bullet m$

processes $P ::= \mathbf{0} \mid \sum_{i \in I} a_i.P_i + \sum_{j \in J} \bar{a}_j \mid \prod_{i \in I} P_i \mid (\text{new } a)P$

r-t processes $R ::= m \triangleright P \mid R \mid R \mid (\text{new } a)R$

transitions $m \triangleright (a.P + Q) \mid m' \triangleright (\bar{a} + R) \longleftrightarrow \langle m', a, Q \rangle \bullet m \triangleright P \mid \langle m, \bar{a}, R \rangle \bullet m' \triangleright \mathbf{0}$

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the encoding of mixed choice in *asynchronous RCCS*

$a.P + \bar{a}$ is encoded as $\llbracket a.P + \bar{a} \rrbracket_c =$

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possible research directions

- coherence is very hard to achieve in nature
 - + biology prompts a thorough study of reversible concurrent calculi where processes have multiplicities and the causal dependencies between copies may be exchanged
- reversible structures may be extended with irreversible combinators (that may be implemented in DNA)
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