A process algebraic approach to reaction systems ☆

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Abstract

In the area of Natural Computing, reaction systems are a qualitative abstraction inspired by the functioning of living cells, suitable to model the main mechanisms of biochemical reactions. This model has already been applied and extended successfully to various areas of research. Reaction systems interact with the environment represented by the context, and pose challenges for further extensions as well as for the implementation, as it is a new computation model. In this paper we consider a variant of the `link`-calculus, which allows to model multiparty interaction in concurrent systems, and show that we can represent reaction systems as processes in a modular way, by representing the behaviour of each entity and preserving faithfully their features. We show the correctness and completeness of our embedding. We illustrate our framework by showing the embedding of a few examples expressing computer science and biological applications. On top of the LTS semantics for reaction systems provided by our framework, we then adapt the classical notion (in concurrency) of bisimulation to make it more suitable for studying properties of reaction systems. In particular, we define a new assertion language based on regular expressions, which allows to specify the properties of interest, and use it to extend Hennessy-Milner logic to our framework. Interestingly, the novel notion of bisimilarity and logical equivalence that are defined parametrically on some assertion of interest are proved to coincide, like in the classical case. Finally, our methodology can contribute to increase the expressiveness of reaction systems, by exploiting the interaction among different reaction systems.

Keywords: process algebras, reaction systems, assertion language, HM-logic

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1. Introduction

Natural Computing is an emerging area of research which has two main aspects: human designed computing inspired by nature, and computation performed in nature. Reaction Systems (RSs) [1] are a rewriting formalism inspired by the way biochemical reactions take place in living cells. This theory has already shown to be relevant in several different fields, such as computer science [2], biology [3, 4, 5, 6], molecular chemistry [7]. Reaction Systems formalise the mechanisms of biochemical systems, such as *facilitation* and *inhibition*. As a qualitative approximation of the real biochemical reactions, they consider if a necessary reagent is or not present, and likewise they consider if an inhibiting molecule is or not present. The possible reactants and inhibitors are called 'entities'. RSs model in a direct way the interaction of a living cell with the environment (called 'context'). However, two RSs are seen as independent models and do not interact.

In this paper, we present an encoding from RSs, to the open multiparty process algebra cCNA,[1] a variant of the `link`-calculus [8, 9] without name mobility. This formalism allows several processes to synchronise and communicate altogether, at the same time, with a new interaction mechanism based on links and link chains. The initial motivation for introducing this mechanism was to encode Mobile Ambients [10], getting a much stronger operational correspondence than any available in the literature, such as the one in [11]. Later it was shown that the `link`-calculus allowed one to easily encode calculi for biology equipped with membranes, as in [12].

We illustrate our embedding by means of some examples coming both from the computer science and the biological field. We also show that our embedding preserves the main features of RSs, and prove its correctness and completeness from the operational semantics viewpoint.

Then, we present a methodology for verifying formally properties of Reaction Systems. The classical notion of bisimulation for process algebras allows to consider two processes as equivalent when one process can simulate all the actions executed by the other one and vice versa. In this paper we define a new notion of bisimilarity which takes into account the characteristics of biological systems. We define a new *bio-simulation* having in mind the possibility that two interacting systems may be compared w.r.t. a subset of the possible biological actions. This is useful for concentrating on the sub-model that one may need to consider for a specific study or application, without getting lost in the complexity of the full biological system, or network. The notion of bio-simulation relies on a new and simple assertion language, which allows to focus on some properties of the Reaction Systems to be verified. In fact, bio-similarity is parametric on a given assertion of interest. Then we prove that the well-known logical characterisation of bisimilarity in terms of Hennessy-Milner Logic [13] (HML) can be extended to the case of bio-similarity by tailoring HML formulas to the same assertion of

[1] After 'chained Core Network Algebra'.

interest used in bio-similarity. As HML is a powerful formalism for specifying properties of labeled transition systems, its variant introduced here is a suitable option to specify formulas over complex labels by abstracting from unnecessary details.

Our main contributions are as follows:

- the behaviour of the context, for each single entity, can be specified in a recursive way, as ordinary processes;
- along the same lines, non-deterministic contexts can be seamlessly expressed in our setting;
- when deterministic contexts are considered, the cCNA computation is deterministic and mimics the evolution of the corresponding RS;
- we define an assertion language to specify local properties of Reaction Systems and exploit it to define a novel notion of behavioural equivalence for Reaction Systems, called bio-similarity;
- we show that our assertions can be used to extend the Hennessy-Milner logic to our framework, in such a way that logical equivalence of processes coincide with bio-similarity;
- we sketch how one can express the behaviour of entity mutation, in such a way that the mutated entity s' can take part to only a subset of rules requiring entity s;
- we show that with a little coding effort, our encoding allows two RSs to communicate; i.e. we can model scenarios where a subset of those entities that the context can provide, are instead provided by a second RS.

The main drawback of our proposal, is that the cCNA translation is verbose. Nevertheless it is clear that our translation can be automatised by means of a proper front-end in an implementation of the `link`-calculus. The examples that we propose also show that some optimisations are possible to reduce the coding when suitable assumptions are made about the provision of entities.

As we have remarked, in our translation, Reaction Systems get the ability to interact between them in a synchronized manner. This interaction is not foreseen in the basic RS framework, as it can only happen with the context. By exploiting recursion, the kind of interactions which can be defined can be complex and expressive. Example 33 and more in general the discussion in Section 7 show that the interaction between RSs can help to model new scenarios.

Related work. Process calculi have been used successfully to model biological processes, see [14] for a recent survey. We are not aware of any SOS operational semantics for the RS; it seems not the case that a deterministic behaviour, as the one of the RS, once the initial state is fixed, could be defined by inference rules that classically are applied to define non-deterministic transition systems. The reversible computation paradigm for RS extends the RS framework by allowing

backward computations as well as forward computations; for this goal a set of inference rules for rewriting logic has been defined in [15] to exactly keep trace of those elements that dissolve in the next computation step.

A preliminary version of this paper appeared in [16]. There are several major differences w.r.t. the paper [16], which is here extended as follows:

- we present several new examples to illustrate our framework, and give more detailed explanations about its use;
- we introduce an assertion language to specify the properties of Reaction Systems;
- we define the notion of bio-simulation to relate Reaction Systems and compare their behaviours;
- we show that our assertion language allows to specify properties extending to our framework the Hennessy-Milner logic;
- we include here all proofs of main results.

Structure of the paper. Section 2 describes RSs and their semantics (interactive processes). Section 3 describes briefly the cCNA process algebra and its operational semantics. Section 4 defines the embedding of RSs in cCNA processes and shows some simple examples to illustrate it. Section 5 presents a couple of examples with computer science and biological applications. Section 6 presents a methodology for the formal verification of properties of Reaction Systems that are expressed in a novel assertion language. Section 7 presents some features and advantages of our embedding for the compositionality of RSs. Finally, Section 8 discusses future work, and concludes.

2. Reaction Systems

Natural Computing is concerned with human-designed computing inspired by nature as well as with computation taking place in nature. The theory of Reaction Systems [1] was born in the field of Natural Computing to model the behaviour of biochemical reactions taking place in living cells. Despite its initial aim, this formalism has shown to be quite useful not only for modeling biological phenomena, but also for the contributions which is giving to computer science [2], theory of computing, mathematics, biology [3, 4, 5, 6], and molecular chemistry [7]. Here we briefly review the basic notions of RSs, see [1] for more details.

The mechanisms that are at the basis of biochemical reactions and thus regulate the functioning of a living cell, are *facilitation* and *inhibition*. These mechanisms are reflected in the basic definitions of Reaction Systems.

Definition 1 (Reaction). A reaction is a triplet $a = (R, I, P)$, where R, I, P are finite, non empty sets and $R \cap I = \emptyset$. If S is a set such that $R, I, P \subseteq S$, then a is a reaction in S.

The sets R, I, P are also written R_a, I_a, P_a and called the *reactant set* of a, the *inhibitor set* of a, and the *product set* of a, respectively. All reactants are needed for the reaction to take place. Any inhibitor blocks the reaction if it is present. Products are the outcome of the reaction. Also, $R_a \cup I_a$ is the set of the resources of a and $rac(S)$ denotes the set of all reactions in S. Because R and I are non empty, all products are produced from at least one reactant and every reaction can be inhibited in some way. Sometimes artificial inhibitors are used that are never produced by any reaction. For the sake of simplicity, in some examples, we will allow I to be empty.

Definition 2 (Reaction System). A Reaction System (RS) is an ordered pair $\mathcal{A} = (S, A)$ such that S is a finite set, and $A \subseteq rac(S)$.

The set S is called the *background set* of $\mathcal{A}$, its elements are called *entities*, they represent molecular substances (e.g., atoms, ions, molecules) that may be present in the states of a biochemical system. The set A is the set of *reactions* of $\mathcal{A}$. Since S is finite, so is A: we denote by $|A|$ the number of reactions in A.

Definition 3 (Reaction Result). Let W be a finite subset of S.

1. Let a be a reaction in S. Then a is enabled by W, denoted by $en_a(W)$, if $R_a \subseteq W$ and $I_a \cap W = \emptyset$. The result of a on W, denoted by $res_a(W)$, is defined by: $res_a(W) = P_a$, if $en_a(W)$, and $res_a(W) = \emptyset$ otherwise.
2. Let A be a finite set of reactions. The result of A on W, denoted by $res_A(W)$, is defined by: $res_A(W) = \bigcup_{a \in A} res_a(W)$.

The theory of Reaction Systems is based on the following assumptions.

- **No permanency.** An entity of a set W vanishes unless it is sustained by a reaction. This reflects the fact that a living cell would die for lack of energy, without chemical reactions.
- **No counting.** The basic model of RSs is very abstract and qualitative, i.e. the quantity of entities that are present in a cell is not taken into account.
- **Threshold nature of resources.** From the previous item, we assume that either an entity is available and there is enough of it (i.e. there are no conflicts), or it is not available at all.

The dynamic behaviour of a RS is formalized in terms of *interactive processes*.

Definition 4 (Interactive Process). Let $\mathcal{A} = (S, A)$ be a RS and let $n \geq 0$. An n-step *interactive process* in $\mathcal{A}$ is a pair $\pi = (\gamma, \delta)$ of finite sequences s.t. $\gamma = \{C_i\}_{i \in [0,n]}$ and $\delta = \{D_i\}_{i \in [0,n]}$ where $C_i, D_i \subseteq S$ for any $i \in [0, n]$, $D_0 = \emptyset$, and $D_i = res_A(D_{i-1} \cup C_{i-1})$ for any $i \in [1, n]$.

Living cells are seen as open systems that continuously react with the external environment, in discrete steps. The sequence γ is the *context sequence* of π and represents the influence of the environment on the Reaction System. The sequence δ is the *result sequence* of π and it is entirely determined by γ and A. The sequence $\tau = W_0, \ldots, W_n$ with $W_i = C_i \cup D_i$, for any $i \in [0, n]$ is called a *state sequence*. Each state W_i in a state sequence is the union of two sets: the context C_i at step i and the result $D_i = res_A(W_{i-1})$ from the previous step.

For technical reasons, we extend the notion of an interactive process to deal with infinite sequences.

Definition 5 (Extended Interactive Process). Let $\mathcal{A} = (S, A)$ be a RS, and let $\pi = (\gamma, \delta)$ be an n-step interactive process, with $\gamma = \{C_i\}_{i\in[0,n]}$ and $\delta = \{D_i\}_{i\in[0,n]}$ Then, we let $\pi^\infty = (\gamma^\infty, \delta^\infty)$ be the extended interactive process of π, defined as $\gamma^\infty = \{C'_i\}_{i\in\mathbb{N}}$, $\delta^\infty = \{D'_i\}_{i\in\mathbb{N}}$, where:

$$C'_j = \begin{cases} C_j & \text{if } j \in [0, n] \\ \emptyset & \text{if } j > n \end{cases} \qquad D'_j = \begin{cases} D_0 & \text{if } j = 0 \\ res_A(D'_{j-1} \cup C'_{j-1}) & \text{if } (j \geq 1) \end{cases}$$

Given an extended interactive process $\pi = (\gamma, \delta)$, we denote by π^k the shift of π starting at the k-th state sequence; formally we let $\pi^k = (\gamma^k, \delta^k)$ with $\gamma^k = \{C'_i\}_{i\in\mathbb{N}}$, $\delta^k = \{D'_i\}_{i\in\mathbb{N}}$ with $C'_0 = C_k \cup D_k$, $D'_0 = \emptyset$, and $C'_i = C_{i+k}$, $D'_i = D_{i+k}$ for any $i \geq 1$.

3. Chained CNA (cCNA)

In this section we introduce the syntax and operational semantics of a variant of the `link`-calculus [8], the cCNA (chained CNA) where the prefixes are link chains and not just links.

Link Chains. Let $\mathcal{C}$ be the set of channels, ranged over by $a, b, ...$, and let $\mathcal{A} = \mathcal{C} \cup \{\tau\} \cup \{\Box\}$ be the set of actions, ranged over by $\alpha, \beta, ...$, where the symbol τ denotes a *silent* action, while the symbol $\Box$ denotes a *virtual* (non-specified) action. A *link* is a pair $\ell = {}^{\alpha}\backslash_{\beta}$; it is *solid* if $\alpha, \beta \neq \Box$; the link ${}^{\Box}\backslash_{\Box}$ is called *virtual*. A link is *valid* if it is solid or virtual. We let $\mathcal{L}$ be the set of valid links. A *link chain* is a finite sequence $v = \ell_1 ... \ell_n$ of (valid) links $\ell_i = {}^{\alpha_i}\backslash_{\beta_i}$ such that:

1. for any $i \in [1, n-1]$, $\begin{cases} \beta_i, \alpha_{i+1} \in \mathcal{C} & \text{implies } \beta_i = \alpha_{i+1} \\ \beta_i = \tau & \text{iff } \alpha_{i+1} = \tau \end{cases}$
2. $\exists i \in [1, n]$. $\ell_i \neq {}^{\Box}\backslash_{\Box}$.

Virtual links represent missing elements of a chain. A chain is called *solid* if it does not contain any virtual link. The equivalence $\bowtie$ models expansion/contraction of virtual links to adjust the length of a link chain.

Definition 6 (Equivalence $\bowtie$). We let $\bowtie$ be the least equivalence relation over link chains closed under the axioms (whenever both sides are well defined):

$$v {}^{\Box}\backslash_{\Box} \bowtie v \qquad {}^{\Box}\backslash_{\Box} v \bowtie v \qquad v_1 {}^{\Box}\backslash_{\Box}\backslash_{\Box} v_2 \bowtie v_1 {}^{\Box}\backslash_{\Box} v_2 \qquad v_1 {}^{\alpha}\backslash_{a}\backslash_{\beta} v_2 \bowtie v_1 {}^{\alpha}\backslash_{a}\backslash_{\Box}\backslash_{\beta} v_2$$

Two link chains of equal length can be merged whenever each position occupied by a solid link in one chain is occupied by a virtual link in the other chain and solid links in adjacent positions match. Positions occupied by virtual links in both chains remain virtual. Merging is denoted by $\upsilon_1 \bullet \upsilon_2$. For example, given $\upsilon_1 = {}^{\tau}\backslash{}^{\Box}_{a}\backslash{}^{\Box}_{\Box}\backslash_{\Box}$, $\upsilon_2 = {}^{\Box}\backslash{}^{a}_{\Box}\backslash{}^{\Box}_{b}\backslash_{\Box}$ and $\upsilon = {}^{\tau}\backslash{}^{a}_{a}\backslash{}^{\Box}_{b}\backslash_{\Box}$ we have $\upsilon_1 \bullet \upsilon_2 = \upsilon$, whereas $\upsilon_1 \bullet \upsilon$ is not defined. Notably the merge operation is commutative and associative.

Some names in a link chain can be restricted as non observable and transformed into silent actions τ. This is possible only if they are matched by some adjacent link. Restriction is denoted by $(\nu\, a)\upsilon$. For example, given $\upsilon = {}^{\tau}\backslash{}^{a}_{a}\backslash{}^{\Box}_{b}\backslash_{\Box}$ as above, we have $(\nu\, a)\upsilon = {}^{\tau}\backslash{}^{\tau}_{\tau}\backslash{}^{\Box}_{b}\backslash_{\Box}$, whereas $(\nu\, b)\upsilon$ is not defined.

Syntax. The set of cCNA processes, denoted as $\mathcal{P}$ and ranged over by P, Q, is defined by the following grammar:

$$P, Q \quad ::= \quad \textstyle\sum_{i \in I} \upsilon_i.P_i \;\mid\; P|Q \;\mid\; (\nu\, a)P \;\mid\; P[\phi] \;\mid\; A$$

where υ_i is a link chain, ϕ is a channel renaming function, and A is a process identifier for which we assume a definition $A \triangleq P$ is available in a given set Δ of (possibly recursive) process definitions. We let $\mathbf{0}$, the inactive process, denote the empty summation.

The syntax of cCNA extends that of CNA [9] by allowing to use link chains as prefixes instead of links. For the rest it features nondeterministic (guarded) choice, parallel composition, restriction, relabelling and possibly recursive definitions. Here we do not consider name mobility, which is present instead in the `link`-calculus.

Semantics. The operational semantics of cCNA is defined in the SOS style by the inference rules in Fig.1. The rules are reminiscent of those for Milner's CCS and they essentially coincide with those of CNA in [9]. The only difference is due to the presence of prefixes that are link chains. Briefly: rule (*Sum*) selects one alternative and puts as label a possible contraction/expansion of the link chain in the selected prefix; rule (*Ide*) selects one transition of the defining process for a constant; rule (*Res*) restricts some names in the label (it cannot be applied when $(\nu\, a)\upsilon$ is not defined); rules (*Lpar*) and (*Rpar*) account for interleaving in parallel composition; rule (*Com*) synchronises interactions (it cannot be applied when $\upsilon \bullet \upsilon'$ is not defined).

Example 7. As some simple examples, consider the recursive process definitions $H \triangleq {}^{a}\backslash_{b}.{}^{a}\backslash_{c}.H$ and $K \triangleq {}^{a}\backslash_{b}.K + {}^{a}\backslash_{c}.K$: the former is some sort of forwarder that alternates between providing a link from a to b and to c; the latter is a nondeterministic forwarder that at each step can link a to b or to c.

Analogously to CNA, the operational semantics of cCNA satisfies the so called Accordion Lemma: whenever $P \xrightarrow{\upsilon} P'$ and $\upsilon' \bowtie \upsilon$ then $P \xrightarrow{\upsilon'} P'$.

$$\frac{\upsilon \bowtie \upsilon_j \quad j \in I}{\sum_{i\in I} \upsilon_i.P_i \xrightarrow{\upsilon} P_j}\ (Sum) \qquad \frac{P \xrightarrow{\upsilon} P' \qquad (A \triangleq P) \in \Delta}{A \xrightarrow{\upsilon} P'}\ (Ide)$$

$$\frac{P \xrightarrow{\upsilon} P'}{(\nu\, a)P \xrightarrow{(\nu\, a)\upsilon} (\nu\, a)P'}\ (Res) \qquad \frac{P \xrightarrow{\upsilon} P'}{P|Q \xrightarrow{\upsilon} P'|Q}\ (Lpar)$$

$$\frac{P \xrightarrow{\upsilon'} P' \qquad Q \xrightarrow{\upsilon} Q'}{P|Q \xrightarrow{\upsilon\bullet\upsilon'} P'|Q'}\ (Com)$$

Figure 1: SOS semantics of the cCNA (rules (*Rel*) and (*Rpar*) omitted).

3.1. Notation for link chains

Hereafter we make use of some new notations for link chains that will facilitate the presentation of our translation.

Definition 8 (Replication). Let υ be a link chain such that $\upsilon\upsilon$ is also a valid link chain. Its n times replication υ^n is defined recursively by letting $\upsilon^0 = \epsilon$ (i.e. the empty chain) and $\upsilon^n = \upsilon\upsilon^{n-1}$.

For example, the expression $({}^{a}\backslash^{\Box}_{b}\backslash_{\Box})^3$ denotes the chain ${}^{a}\backslash^{\Box}_{b}\backslash^{a}_{\Box}\backslash^{\Box}_{b}\backslash^{a}_{\Box}\backslash^{\Box}_{b}\backslash_{\Box}$. Instead the expression $({}^{a}\backslash_{b})^2$ is ill-defined because a does not match with b.

Then, we introduce the notation for *half links* that will be used in conjunction with the *open block of chain* to form regular link chains.

Definition 9 (Half links). Let a be a channel name, we define the *half left link* ${}^{a}\backslash$, and the *half right link* $\backslash_a$.

Definition 10 (Open block). Let R be a set of names. We define an *open block* as $\left(\Vert_{a\,\in\,R} {}^{\Box}_{a_i}\backslash^{a_o}_{\Box}\right)$, where a_i and a_o are annotated version of the name a, as

set	block of chain	result
$R = \emptyset$	$\left(\Vert_{a\,\in\,R} {}^{\Box}_{a_i}\backslash^{a_o}_{\Box}\right)$	ϵ
$R = \{b\}$	$\left(\Vert_{a\,\in\,R} {}^{\Box}_{a_i}\backslash^{a_o}_{\Box}\right)$	${}^{\Box}_{b_i}\backslash^{b_o}_{\Box}$
$R = \{b\} \cup R'$	$\left(\Vert_{a\,\in\,R} {}^{\Box}_{a_i}\backslash^{a_o}_{\Box}\right)$	${}^{\Box}_{b_i}\backslash^{b_o}_{\Box} \backslash \left(\Vert_{a\,\in\,R'} {}^{\Box}_{a_i}\backslash^{a_o}_{\Box}\right)$

We then combine half links and open blocks to form regular link chains. For example, for $R = \{a, b\}$ the expression $\left(\Vert_{c\,\in\,R} {}^{\Box}_{c_i}\backslash^{c_o}_{\Box}\right)$ denotes the block ${}^{\Box}_{a_i}\backslash^{a_o}_{\Box}\backslash^{\Box}_{b_i}\backslash^{b_o}_{\Box}$; and the expression ${}^{r_1}\backslash \left(\Vert_{c\,\in\,R} {}^{\Box}_{c_i}\backslash^{c_o}_{\Box}\right) \backslash_{r_2}$ denotes the chain ${}^{r_1}\backslash^{\Box}_{a_i}\backslash^{a_o}_{\Box}\backslash^{\Box}_{b_i}\backslash^{b_o}_{\Box}\backslash_{r_2}$.

4. From Reaction Systems to cCNA

Here we present a translation from Reaction Systems to cCNA. The idea is to define separated processes for representing the behaviour of each entity, each reaction, and for the provisioning of each entity by the context.

Processes for entities. Given an entity $s \in S$, we exploit five different pairs of names for the interactions over s: names s_i, s_o are used to test the presence of s in the system; names $\widehat{s}_i$, $\widehat{s}_o$ are used to test the provisioning of s from the context; names $\widetilde{s}_i$, $\widetilde{s}_o$ are used to test the production of s by some reaction; names $\overline{s}_i$, $\overline{s}_o$ are used to test the absence of s in the system; and names $\underline{s}_i$, $\underline{s}_o$ are used to test the absence of s from the context. We let P_s be the process implementing the presence of s in the system, and $\overline{P_s}$ be the one for its absence. They can be seen as instances of the same template, which is given below.

$$\begin{array}{rcl} P_s & \triangleq & E(s,\widetilde{s},\widehat{s},\underline{s}) \qquad\qquad \overline{P_s} \triangleq E(\overline{s},\widetilde{s},\widehat{s},\underline{s}) \\ E(s,\widetilde{s},\widehat{s},\underline{s}) & \triangleq & \sum_{h,k\geq 0}({}^{s_i}\backslash^{\square}_{s_o}\backslash\square)^h\ {}^{\widehat{s}_i}\backslash^{\square}_{\widehat{s}_o}\backslash\square({}^{\widetilde{s}_i}\backslash^{\square}_{\widetilde{s}_o}\backslash\square)^k.P_s \\ & & + \\ & & \sum_{h\geq 0,k\geq 1}({}^{s_i}\backslash^{\square}_{s_o}\backslash\square)^h\ {}^{\underline{s}_i}\backslash^{\square}_{\underline{s}_o}\backslash\square({}^{\widetilde{s}_i}\backslash^{\square}_{\widetilde{s}_o}\backslash\square)^k.P_s \\ & & + \\ & & \sum_{h\geq 0}({}^{s_i}\backslash^{\square}_{s_o}\backslash\square)^h\ {}^{\underline{s}_i}\backslash_{\underline{s}_o}.\overline{P_s} \end{array}$$

The first line of $E(s,\widetilde{s},\widehat{s},\underline{s})$ accounts for the case where s is tested for presence by h reactions and produced by k reactions, while being provided by the context (${}^{\widehat{s}_i}\backslash_{\widehat{s}_o}$). Thus, s will be present at the next step (the continuation is P_s). Here h and k are not known a priori and therefore any combination is possible. In practice, by knowing the number of reactions that test s, we can bound the maximum values of h and k. The second line accounts for the analogous case where s is not provided by the context (${}^{\underline{s}_i}\backslash_{\underline{s}_o}$). The condition $k \geq 1$ guarantees that s will remain present (the continuation is P_s). The third line accounts for the case where s is tested for presence, but it is neither produced nor provided by the context. Therefore, in the next step s will be absent in the system (the continuation is $\overline{P_s}$). Note that in the case of $\overline{P_s}$ the test for presence of s in the system is just replaced by the test for its absence.

Processes for reactions. We assume that each reaction a is assigned a progressive number j. The process for reaction $aj = (R_j, I_j, P_j)$ must assert either the possibility to apply the reaction or its impossibility. The first case happens when all its reactants are present (the link ${}^{s_i}\backslash_{s_o}$ is requested for any $s \in R_j$) and all its inhibitors are absent (the link ${}^{\overline{e}_i}\backslash_{\overline{e}_o}$ is requested for any $e \in I_j$), then the product set is released (the link ${}^{\widetilde{c}_i}\backslash_{\widetilde{c}_o}$ is requested for any $c \in P_j$). The next case can happen for two reasons: one of the reactants is absent (the link ${}^{\overline{s}_i}\backslash_{\overline{s}_o}$ is requested for some $s \in R_j$) or one of the inhibitors is present (the link ${}^{e_i}\backslash_{e_o}$ is requested for some $e \in I_j$). The process is recursive so that reactions can be

applied at any step.

$$
\begin{array}{ll}
P_{aj} \triangleq & \\
{}^{r_j}\backslash \left(\Vert_{s \in R_j} \overset{\square}{s_i} \backslash \overset{s_o}{\square} \right) \backslash \left(\Vert_{e \in I_j} \overset{\square}{\overline{e}_i} \backslash \overset{\overline{e}_o}{\square} \right) \backslash {}^{\square}_{r_{j+1}} \backslash {}^{p_j}_{\square} \backslash \left(\Vert_{c \in P_j} \overset{\square}{\widetilde{c}_i} \backslash \overset{\widetilde{c}_o}{\square} \right) \backslash_{p_{j+1}} . P_{aj} & \{aj \text{ is applicable}\} \\
+ & \\
\sum_{s \in R_j} {}^{r_j}\backslash \overset{\square}{\overline{s}_i} \backslash \overset{\overline{s}_o}{\square} \backslash {}^{\square}_{r_{j+1}} \backslash {}^{p_j}_{\square} \backslash_{p_{j+1}} . P_{aj} & \{aj \text{ is not applicable}\} \\
+ & \\
\sum_{e \in I_j} {}^{r_j}\backslash \overset{\square}{e_i} \backslash \overset{e_o}{\square} \backslash {}^{\square}_{r_{i+1}} \backslash {}^{p_j}_{\square} \backslash_{p_{j+1}} . P_{aj} & \{aj \text{ is not applicable}\}
\end{array}
$$

We exploit names r_j, p_j to join the chains provided by the application of all the reactions. Channels r_j and r_{j+1} enclose the enabling/disabling condition of reaction aj. Channels p_j and p_{j+1} enclose the links related to the entities produced by aj. We will see that all the link chain labels of transitions follow the same schema: first we find all the reactions limited to the reactants and inhibitors (chained using r_j channels), then all the supplies by the contexts (chained using the channel *cxt*, to be introduced next), and finally the products for all the reactions (chained using p_j channels). In the following there is an example explaining this schema.

Processes for contexts. The context can vary at each step n and for each entity $s \in S$, the context must say if the entity is provided or not. Let us denote by C_n the context at step n, i.e. the set of entities that are made available. Correspondingly, we introduce another process Cxt^n defined as follows:

$$
Cxt^n \quad \triangleq \quad {}^{cxt}\backslash \left(\Vert_{s \in C_n} \overset{\square}{\hat{s}_i} \backslash \overset{\hat{s}_o}{\square} \right) \backslash \left(\Vert_{e \notin C_n} \overset{\square}{\underline{e}_i} \backslash \overset{\underline{e}_o}{\square} \right) \backslash_{p_1} . Cxt^{n+1}
$$

We only consider Cxt^n with $n > 0$, as the entities that are present at step zero are considered to be present in the initial system (if $s \in C_0$ the process P_s will be present initially, otherwise $\overline{P_s}$ will be present).

In the following we use the following conventions for denoting different categories of names:

- *decs* $\triangleq \{s, \overline{s}, \widetilde{s}, \widehat{s}, \underline{s} \mid s \in S\}$ is the set of names for decorated entities (without subscripts $_i$ and $_o$);
- *ents* $\triangleq \{d_i, d_o \mid d \in decs\}$ is the set names for entities;
- *reacts* $\triangleq \{r_1, ..., r_{|A|}\}$ is the set of reaction names r_j associated with each reaction aj;
- *prods* $\triangleq \{p_1, ..., p_{|A|}\}$ is the set of names for production p_j, associated with reaction aj.

Definition 11 (Translation). Let $\mathcal{A} = (S, A)$ be a RS, and let $\pi = (\gamma, \delta)$ be an extended interactive process in $\mathcal{A}$, with $\gamma = \{C_i\}_{i \in \mathbb{N}}$. We define its

cCNA translation $[\![\mathcal{A}, \gamma]\!]$ as follows:

$$[\![\mathcal{A}, \gamma]\!] \triangleq (\nu\, names)\left(I \mid \prod_{a \in A} P_a \mid Cxt^1 \mid \prod_{s \in C_0} P_s \mid \prod_{s \notin C_0} \overline{P_s}\right)$$

where $names = reacts \cup ents \cup prods \cup \{cxt\}$. For technical reasons, we introduce the init process $I \triangleq {}^{\tau}\backslash_{r_1}.I$ to allow the name r_1 to be matched at the start of any chain; also, for notational convenience, we fix that $r_{u+1} = cxt$ and $p_{u+1} = \tau$, for $u = |A|$.

It is important to observe that, for each transition, our cCNA encoding requires all the processes running in parallel to interact in that transition. This is due to the fact that all the channels r_j, p_j, cxt, including those for decorated names $s_i, s_o, \overline{s}_i, \overline{s}_o$ are restricted. Each reaction defines a pattern to be satisfied, i.e. each reaction inserts as many virtual links as the number of reactants, inhibitors, and products, as required by the corresponding reaction.

Lemma 12. *Let $\mathcal{A} = (S, A)$ be a RS and let $\pi = (\gamma, \delta)$ be an extended interactive process in $\mathcal{A}$. Let $P = [\![\mathcal{A}, \gamma]\!]$ its cCNA translation. If exists P' such that $P \xrightarrow{(\nu\, names)\upsilon} P'$ is a transition of P, then*

1. *for each reaction $aj \in A$, the corresponding channels r_j and p_j appear in υ; for each entity $s \in S$, the corresponding channel s (suitably decorated) appear in υ; the channel cxt appears in υ;*
2. *for each reaction $aj \in A$ and each entity $s \in S$, each virtual link offered by processes P_a and Cxt is overlapped by exactly one solid link offered by processes representing entities.*

The topmost restriction $(\nu\, names)$ appearing in the process $[\![\mathcal{A}, \gamma]\!]$ serves to guarantee that all names appearing in a link of the chain labelling a transition are matched. Since all names appearing in any prefix of $[\![\mathcal{A}, \gamma]\!]$ are restricted, in the transition $[\![\mathcal{A}, \gamma]\!] \xrightarrow{(\nu\, names)\upsilon} P'$ it means that the observation $(\nu\, names)\upsilon$ has the form ${}^{\tau}\backslash_{\tau} \ldots {}^{\tau}\backslash_{\tau}$, i.e., it is silent, and that υ is solid. Later on we will be interested in reasoning about the actual chain υ used in the transition. It has the peculiarity to start and end with silent actions and to include all names in $reacts \cup prods \cup \{cxt\}$. As a matter of notation we call such chain υ *complete*.

Definition 13 (Complete Chain). A chain υ is called *complete* if it is solid (i.e., it contains no virtual link) and has silent actions τ at its extremes. We write $P \overset{\upsilon}{\Rightarrow} P'$ to mean that $P \xrightarrow{\upsilon} P'$ with υ complete.

We will then use $(\!|\mathcal{A}, \gamma|\!)$ to refer to the encoding without topmost name restrictions, i.e.,

$$(\!|\mathcal{A}, \gamma|\!) \triangleq I \mid \prod_{a \in A} P_a \mid Cxt^1 \mid \prod_{s \in C_0} P_s \mid \prod_{s \notin C_0} \overline{P_s}.$$

and we will focus on the complete transitions of $(\!|\mathcal{A}, \gamma|\!)$.

The following Corollary immediately follows from Lemma 12.

Corollary 14. *Let $\mathcal{A} = (S, A)$ be a RS and let $\pi = (\gamma, \delta)$ be an extended interactive process in $\mathcal{A}$. Let $Q = (\!|\mathcal{A}, \gamma|\!)$ and $P = [\![\mathcal{A}, \gamma]\!] = (\nu\, names)Q$. Then $P \xrightarrow{(\nu\, names)\upsilon} P'$ iff $Q \overset{\upsilon}{\Rightarrow} Q'$ and $P' = (\nu\, names)Q'$.*

Example 15. Let $\mathcal{A}$ be a RS whose specification contains two entities, $s1$ and $s2$, and the reactions $r_1 = (s1, \ , s2)$ and $r_2 = (s2, \ , s1)$ that produce $s2$ if $s1$ is present and $s1$ if $s2$ is present. For simplicity we do not consider inhibitors, but the reader can assume a void inhibitor is present in both reactions. Then, we assume an extended interactive process $\pi = (\gamma, \delta)$ where the context γ provides $s1$ and $s2$ at every step, but we assume that only $s1$ is initially present. The corresponding cCNA process is $[\![\mathcal{A}, \gamma]\!] \triangleq (\nu\, names)(\!|\mathcal{A}, \gamma|\!)$, with

$$(\!|\mathcal{A}, \gamma|\!) \triangleq I \mid P_{s1} \mid \overline{P}_{s2} \mid P_{r1} \mid P_{r2} \mid Cxt$$

where:

$$\begin{aligned}
P_{r_1} &\triangleq r_1 \backslash^{\square}_{s1_i} \backslash^{s1_o}_{\square} \backslash^{\square}_{r_2} \backslash^{p_1}_{\square} \backslash^{\square}_{\widetilde{s2}_i} \backslash^{\widetilde{s2}_o}_{\square} \backslash_{p_2}.P_{r_1} + \ldots; \\
P_{r_2} &\triangleq \cdots + r_2 \backslash^{\square}_{\widehat{s2}_i} \backslash^{\overline{s2}_o}_{\square} \backslash^{\square}_{cxt} \backslash^{p_2}_{\square} \backslash_{\tau}.P_{r_2} + \ldots; \\
P_{s1} &\triangleq s1_i \backslash^{\square}_{s1_o} \backslash^{\widehat{s1}_i}_{\square} \backslash_{\widehat{s1}_o}.P_{s1} + \ldots; \\
\overline{P}_{s2} &\triangleq \overline{s2}_i \backslash^{\square}_{\overline{s2}_o} \backslash^{\widehat{s2}_i}_{\square} \backslash^{\square}_{\widehat{s2}_o} \backslash^{\widetilde{s2}_i}_{\square} \backslash_{\widetilde{s2}_o}.P_{s2} + \ldots; \\
Cxt &\triangleq cxt \backslash^{\square}_{\widehat{s1}_i} \backslash^{\widehat{s1}_o}_{\square} \backslash^{\square}_{\widehat{s2}_i} \backslash^{\widehat{s2}_o}_{\square} \backslash_{p_1}.Cxt
\end{aligned}$$

For clarity of exposition, we show the code of the processes just in part, to focus on the prefixes that will be involved in the first transition of the system. In Figure 2 we show the structure of a link chain label related to the execution of such a transition. The yellow blocks are referred to init process I, to the processes encoding the reactions, P_{r_1} and P_{r_2}, and to the context Cxt. As the figure puts in evidence, these two kinds of processes determine the structure of the link chain, from end to end, i.e. from the left τ to the right one. We could say that these processes form the *backbone* of the interaction. In contrast, the processes encoding the entities, P_{s1}, $\overline{P}_{s2}$, provide the solid links to be merged with the virtual links of the backbone (i.e. to be plugged in the backbone). In Figure 2 we added the underline brackets, where the notation $P_1(P_2, P_3)$ means that the segment of the link chain is delimited by the process P_1 and it leaves "holes" where processes between the brackets insert their links. Formally, we have

$$(\!|\mathcal{A}, \gamma|\!) \xrightarrow{\tau \backslash^{r_1}_{r_1} \backslash^{s1_i}_{s1_i} \backslash^{s1_o}_{s1_o} \backslash^{r_2}_{r_2} \backslash^{\overline{s2}_i}_{\overline{s2}_i} \backslash^{\overline{s2}_o}_{\overline{s2}_o} \backslash^{cxt}_{cxt} \backslash^{\widehat{s1}_i}_{\widehat{s1}_i} \backslash^{\widehat{s1}_o}_{\widehat{s1}_o} \backslash^{\widehat{s2}_i}_{\widehat{s2}_i} \backslash^{\widehat{s2}_o}_{\widehat{s2}_o} \backslash^{p_1}_{p_1} \backslash^{\widetilde{s2}_i}_{\widetilde{s2}_i} \backslash^{\widetilde{s2}_o}_{\widetilde{s2}_o} \backslash^{p_2}_{p_2} \backslash_{\tau}} P'$$

Since the chain in the transition label is complete we can also write

$$(\!|\mathcal{A}, \gamma|\!) \xRightarrow{\tau \backslash^{r_1}_{r_1} \backslash^{s1_i}_{s1_i} \backslash^{s1_o}_{s1_o} \backslash^{r_2}_{r_2} \backslash^{\overline{s2}_i}_{\overline{s2}_i} \backslash^{\overline{s2}_o}_{\overline{s2}_o} \backslash^{cxt}_{cxt} \backslash^{\widehat{s1}_i}_{\widehat{s1}_i} \backslash^{\widehat{s1}_o}_{\widehat{s1}_o} \backslash^{\widehat{s2}_i}_{\widehat{s2}_i} \backslash^{\widehat{s2}_o}_{\widehat{s2}_o} \backslash^{p_1}_{p_1} \backslash^{\widetilde{s2}_i}_{\widetilde{s2}_i} \backslash^{\widetilde{s2}_o}_{\widetilde{s2}_o} \backslash^{p_2}_{p_2} \backslash_{\tau}} P'$$

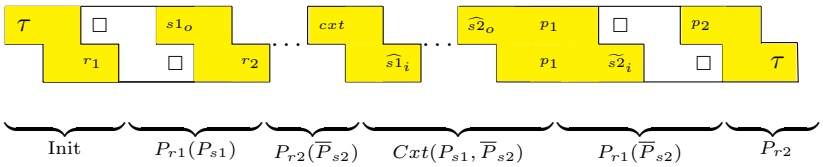

Figure 2: The link chain structure arising from reactions and context processes.

Example 15 outlines two different roles of the processes defining the translation of an interactive process: those processes encoding the reactions and the context provide the backbone of each transition, whereas the processes encoding the entities provide the resources needed for the communication to take place.

The flat function. For technical reasons, our transition labels are quite verbose; then, to simplify their processing, we introduce a function that takes a solid link chain and returns a simple string by eliminating all the channel matching pairs leaving just one placeholder for them. This transformation is harmless, in the sense that it retains all the information in the chain, because it is applied to complete chains only. The function $\mathit{flat}(\cdot)$ is defined inductively as follows:

$$\mathit{flat}(\epsilon) \triangleq \epsilon \qquad \mathit{flat}({}^{\alpha}\backslash_{\beta}) \triangleq \begin{cases} \beta & \text{if } \beta \in \mathit{reacts} \cup \{\mathit{cxt}\} \cup \mathit{prods} \\ d & \text{if } \beta = d_i \text{ with } d \in \mathit{decs} \\ \epsilon & \text{otherwise} \end{cases}$$

$$\mathit{flat}({}^{\alpha}\backslash_{\beta}\upsilon) \quad \triangleq \quad \mathit{flat}({}^{\alpha}\backslash_{\beta}) :: \mathit{flat}(\upsilon)$$

where :: is the usual string concatenation operator.

For example, if we consider again the complete label

$$\upsilon = {}^{\tau}\backslash^{r_1}_{r_1}\backslash^{s1_i}_{s1_i}\backslash^{s1_o}_{s1_o}\backslash^{r_2}_{r_2}\backslash^{\overline{s2}_i}_{\overline{s2}_i}\backslash^{\overline{s2}_o}_{\overline{s2}_o}\backslash^{\mathit{cxt}}_{\mathit{cxt}}\backslash^{\widehat{s1}_i}_{\widehat{s1}_i}\backslash^{\widehat{s1}_o}_{\widehat{s1}_o}\backslash^{\widehat{s2}_i}_{\widehat{s2}_i}\backslash^{\widehat{s2}_o}_{\widehat{s2}_o}\backslash^{p_1}_{p_1}\backslash^{\widetilde{s2}_i}_{\widetilde{s2}_i}\backslash^{\widetilde{s2}_o}_{\widetilde{s2}_o}\backslash^{p_2}_{p_2}\backslash_{\tau}$$

from Example 15, we have

$$\mathit{flat}(\upsilon) = r_1\ s1\ r_2\ \overline{s2}\ \mathit{cxt}\ \widehat{s1}\ \widehat{s2}\ p_1\ \widetilde{s2}\ p_2.$$

It is then immediate to define the function *unflat* to rebuild the complete label from the compact string (here we exploit again the half link and block notation):

$$\mathit{unflat}(x) \triangleq \begin{cases} x & \text{if } x \in \mathit{reacts} \cup \{\mathit{cxt}\} \cup \mathit{prods} \\ {}^{x}\backslash_{x} & \text{if } x \in \mathit{decs} \end{cases}$$

$$\mathit{unflat}(x_1 \ldots x_n) \triangleq {}^{\tau}\backslash \mathit{unflat}(x_1) \backslash \ldots \backslash \mathit{unflat}(x_n) \backslash_{\tau}$$

It is immediate to check that for any complete label υ of our processes we have $\upsilon = \mathit{unflat}(\mathit{flat}(\upsilon))$.

With the next proposition, we analyse the structure of a cCNA process encoding of a reactive process after one transition step. In the following four statements, for brevity, we let $\mathcal{A} = (S, A)$ be a RS, and let $\pi = (\gamma, \delta)$ be an extended interactive process in A, with $\gamma = \{C_i\}_{i \in \mathbb{N}}$ and $\delta = \{D_i\}_{i \in \mathbb{N}}$.

Proposition 16 (Correctness 1). *Let* $P = [\![\mathcal{A}, \gamma]\!]$ *with*

$$P = (\nu\, names)\left(I \mid \prod_{a\in A} P_a \mid Cxt^1 \mid \prod_{s\in C_0} P_s \mid \prod_{s\notin C_0} \overline{P}_s\right).$$

If there exists P' *such that* $P \xrightarrow{v} P'$*, it holds that:*

1. $v = {}^{\tau}\backslash_{\tau} \ldots {}^{\tau}\backslash_{\tau}$*, and*
2. $P' = (\nu\, names)\left(I \mid \prod_{a\in A} P_a \mid Cxt^2 \mid \prod_{s\in C_1\cup D_1} P_s | \prod_{s\notin C_1\cup D_1} \overline{P}_s\right)$.

Moreover, given $\pi^1 = (\gamma^1, \delta^1)$*, we have* $P' = [\![\mathcal{A}, \gamma^1]\!]$.

Now, we extend the previous result to a series of transitions.

Corollary 17 (Correctness 2). *Let* $P = [\![\mathcal{A}, \gamma]\!]$ *and* $j \geq 1$*. If there exists* P'' *such that* $P \xrightarrow{{}^{\tau}\backslash_{\tau}\ldots{}^{\tau}\backslash_{\tau}}{}^{j} P''$*, then letting* $\pi^j = (\gamma^j, \delta^j)$ *we have* $P'' = [\![\mathcal{A}, \gamma^j]\!]$.

With the following propositions, we prove that, given a RS $\mathcal{A} = (S, A)$ and an extended interactive process $\pi = (\gamma, \delta)$, then the *c*CNA process $[\![\mathcal{A}, \gamma]\!]$ can simulate all the evolutions of π.

Proposition 18 (Completeness 1). *Let* $P = [\![\mathcal{A}, \gamma]\!]$ *and* $\pi^1 = (\gamma^1, \delta^1)$*. Then,* $P \xrightarrow{{}^{\tau}\backslash_{\tau}\ldots{}^{\tau}\backslash_{\tau}} P' = [\![\mathcal{A}, \gamma^1]\!]$.

Now, we extend the previous result to a series of transitions.

Corollary 19 (Completeness 2). *Let* $P = [\![\mathcal{A}, \gamma]\!]$ *and* $\pi^j = (\gamma^j, \delta^j)$*. Then,* $P \xrightarrow{{}^{\tau}\backslash_{\tau}\ldots{}^{\tau}\backslash_{\tau}}{}^{j} P'' = [\![\mathcal{A}, \gamma^j]\!]$.

5. Examples

Hereafter, we shall often omit topmost restrictions but shall take into account only transitions whose labels are complete chains, i.e. they do not contain virtual links and start/end with the τ action symbol.

5.1. Labelled transition system

This example is inspired by the example in [1], where a deterministic transition system is coded in the Reaction System framework. Here we consider the minimal deterministic transition system in Figure 3. In our *c*CNA encoding, q, w, a, and b became the *entities*, the context can only provide a and b and the reaction rules are as follows:

$$\begin{array}{llll} 1 & (\{q,a\},\{w,b\},\{w\}) & 2 & (\{q,b\},\{w,a\},\{q\}) \\ 3 & (\{w,a\},\{q,b\},\{w\}) & 4 & (\{w,b\},\{q,a\},\{q\}) \end{array}$$

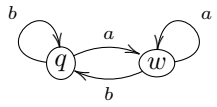

Figure 3: Minimal deterministic labelled transition system.

Encoding of the rules. The encoding of the rules for reactions is given in a parametric way:

$$\begin{array}{lcl} P_n(q,b,w,a,q) & \triangleq & \pi_n(q,b,w,a,q).P_n(q,b,w,a,q) \\ & & + \\ & & \sum_{x\in\{\overline{q},\overline{b},w,a\}}\ \pi'_n(x).P_n(q,b,w,a,q) \end{array}$$

where

$$\begin{array}{lcl} \pi_n(q,b,w,a,q) & \triangleq & {}^{r_n}\backslash^{\Box}_{q_i}\backslash^{q_o}_{\Box}\backslash^{\Box}_{b_i}\backslash^{b_o}_{\Box}\backslash^{\Box}_{\overline{w}_i}\backslash^{\overline{w}_o}_{\Box}\backslash^{\Box}_{\overline{a}_i}\backslash^{\overline{a}_o}_{\Box}\backslash^{\Box}_{r_{n+1}}\backslash^{p_n}_{\Box}\backslash^{\Box}_{\widetilde{q}_i}\backslash^{\widetilde{q}_o}_{\Box}\backslash_{p_{n+1}} \\ \pi'_n(x) & \triangleq & {}^{r_n}\backslash^{\Box}_{x_i}\backslash^{x_o}_{\Box}\backslash^{\Box}_{r_{n+1}}\backslash^{p_n}_{\Box}\backslash_{p_{n+1}} \end{array}$$

Then we have

$$\begin{array}{lcllcl} P_1 & \triangleq & P_1(q,a,w,b,w) & \qquad P_3 & \triangleq & P_3(w,a,q,b,w) \\ P_2 & \triangleq & P_2(q,b,w,a,q) & \qquad P_4 & \triangleq & P_4(w,b,q,a,q) \end{array}$$

and we put, as usual, $r_5 = cxt$ and $p_5 = \tau$.

Encoding of the entities. As for reactions, also the encoding of the entities is given in a parametric way. Here we differentiate the encoding for the entities that are not provided by the context and that can be produced by the reactions, and the ones that can be provided by the context and that are not produced by the reactions.

Here, for the entities q and w that are not provided by the context, we let:

$$\begin{array}{lcllcl} P_q & \triangleq & E(q,\widetilde{q}) & \qquad \overline{P}_q & \triangleq & E(\overline{q},\widetilde{q}) \\ P_w & \triangleq & E(w,\widetilde{w}) & \qquad \overline{P}_w & \triangleq & E(\overline{w},\widetilde{w}) \end{array}$$

where:

$$\begin{array}{lcl} E(q,\widetilde{q}) & \triangleq & \sum_{h=1}^{3}({}^{q_i}\backslash^{\Box}_{q_o}\backslash_{\Box})^h\ {}^{\widetilde{q}_i}\backslash_{\widetilde{q}_o}.P_q \\ & & + \\ & & \sum_{h=1}^{3}({}^{q_i}\backslash^{\Box}_{q_o}\backslash_{\Box})^h.\overline{P}_q \end{array}$$

In fact the presence/absence of q and w will be exploited by at least one rule and at most three rules.

Here, for the entities a and b that can be provided by the context but not produced by the rules, we let:

$$\begin{array}{lcllcl} P_a & \triangleq & E(a,\widehat{a},\underline{a}) & \qquad \overline{P}_a & \triangleq & E(\overline{a},\widehat{a},\underline{a}) \\ P_b & \triangleq & E(b,\widehat{b},\underline{b}) & \qquad \overline{P}_b & \triangleq & E(\overline{b},\widehat{b},\underline{b}) \end{array}$$

where:

$$E(a,\widehat{a},\underline{a}) \triangleq \sum_{h=1}^{3}({}^{a_i}\backslash^{\square}_{a_o}\backslash_{\square})^h\ {}^{\widehat{a}_i}\backslash_{\widehat{a}_o}.P_a + \sum_{h=1}^{3}({}^{a_i}\backslash^{\square}_{a_o}\backslash_{\square})^h\ {}^{\underline{a}_i}\backslash_{\underline{a}_o}.\overline{P}_a$$

Finally, for the context, the encoding follows:

$$Cxt \triangleq {}^{cxt}\backslash^{\square}_{\widehat{a}_i}\backslash^{\widehat{a}_o}_{\square}\backslash^{\square}_{\underline{b}_i}\backslash^{\underline{b}_o}_{\square}\backslash_{p_1}.Cxt \quad + \quad {}^{cxt}\backslash^{\square}_{\widehat{b}_i}\backslash^{\widehat{b}_o}_{\square}\backslash^{\square}_{\underline{a}_i}\backslash^{\underline{a}_o}_{\square}\backslash_{p_1}.Cxt$$

Notice that we exploit here the capabilities of the process algebraic framework to define a nondeterministic, recursive context. We model the context to always offer either a or b, but never both the entities together. The reason is that in the other cases (providing both a and b or neither of them) would lead the system to be stuck because of the simplifications we have adopted in the other processes.

Now, we assume that we have an initial configuration containing the entities q and b:

$$Sys \triangleq I \mid P_q \mid \overline{P}_w \mid \overline{P}_a \mid P_b \mid P_1 \mid P_2 \mid P_3 \mid P_4 \mid Cxt.$$

Then, only the second reaction can be applied, and the transition carries the complete label υ below

$${}^{\tau}\backslash^{r_1}_{r_1}\backslash^{\overline{\mathbf{a}}_\mathbf{i}}_{\overline{a}_i}\backslash^{\overline{a}_o}_{\overline{\mathbf{a}}_\mathbf{o}}\backslash^{r_2}_{r_2}\backslash^{\mathbf{q_i}}_{q_i}\backslash^{q_o}_{\mathbf{q_o}}\backslash^{\mathbf{b_i}}_{b_i}\backslash^{b_o}_{\mathbf{b_o}}\backslash^{\overline{\mathbf{w}}_\mathbf{i}}_{\overline{w}_i}\backslash^{\overline{w}_o}_{\overline{\mathbf{w}}_\mathbf{o}}\backslash^{\overline{\mathbf{a}}_\mathbf{i}}_{\overline{a}_i}\backslash^{\overline{a}_o}_{\overline{\mathbf{a}}_\mathbf{o}}\backslash^{r_3}_{r_3}\backslash^{\overline{\mathbf{a}}_\mathbf{i}}_{\overline{a}_i}\backslash^{\overline{a}_o}_{\overline{\mathbf{a}}_\mathbf{o}}\backslash^{r_4}_{r_4}\backslash^{\overline{\mathbf{w}}_\mathbf{i}}_{\overline{w}_i}\backslash^{\overline{w}_o}_{\overline{\mathbf{w}}_\mathbf{o}}\backslash^{cxt}_{cxt}\backslash^{\widehat{\mathbf{a}}_\mathbf{i}}_{\widehat{a}_i}\backslash^{\widehat{a}_o}_{\widehat{\mathbf{a}}_\mathbf{o}}\backslash^{\underline{\mathbf{b}}_\mathbf{i}}_{\underline{b}_i}\backslash^{\underline{b}_o}_{\underline{\mathbf{b}}_\mathbf{o}}\backslash^{p_1}_{p_1}\backslash^{p_2}_{p_2}\backslash^{\widetilde{\mathbf{q}}_\mathbf{i}}_{\widetilde{q}_i}\backslash^{\widetilde{q}_o}_{\widetilde{\mathbf{q}}_\mathbf{o}}\backslash^{p_3}_{p_3}\backslash^{p_4}_{p_4}\backslash_{\tau}$$

The parts in bold are provided by the entity processes, the other parts are provided by the processes encoding the reactions and by the process encoding the context (starting at cxt and ended at p_1. In the label we can read that the rules 1 and 4 have been not executed because the entity a is absent, the rule 3 has been not applied because the entity w is absent, then only rule 2 has been applied, and it has produced the entity q. Also, the context provides entity a, that will be available in the next state, and not the entity b. Now, to let the label more readable, we show the result of the application of the function $flat(\cdot)$ to it:

$$r_1\ \overline{a}\ r_2\ q\ b\ \overline{w}\ \overline{a}\ r_3\ \overline{a}\ r_4\ \overline{w}\ cxt\ \widehat{a}\ \underline{b}\ p_1\ p_2\ \widetilde{q}\ p_3\ p_4.$$

5.2. A biological toy example of gene expression

We consider a biological toy example in the style of gene's alternative splicing [17]. Alternative splicing is a regulated process during gene expression that results in a single gene coding for multiple proteins. In practice, particular exons of a gene may be included within or excluded from the final, processed messenger RNA (mRNA) produced from that gene. In our example, a gene a codes for a protein T when molecules G is present and C is absent, and in the opposite situation a codes for protein T'. This behavior is encoded in rules 1 and 2. Then, rule 3 codes for the production of C when proteins T and F are present, and T' absent; rule 4 codes for the production of G when proteins T' is present and F is absent.

Encoding of the rules. The encoding of the rules for reactions is given in a parametric way:

$$\begin{array}{lcl} P_n(a,G,C,T) & \triangleq & \pi_n(a,G,C,T).P_n(a,G,C,T) \\ & & + \\ & & \sum_{x\in\{a,G,C\}} \pi'_n(x).P_n(a,G,C,T) \end{array}$$

where

$$\begin{array}{lcl} \pi_n(a,G,C,T) & \triangleq & r_n\backslash^{\square}_{a_i}\backslash^{a_o}_{\square}\backslash^{\square}_{G_i}\backslash^{G_o}_{\square}\backslash^{\square}_{\overline{C}_i}\backslash^{\overline{C}_o}_{\square}\backslash^{\square}_{r_{n+1}}\backslash^{p_n}_{\square}\backslash^{\square}_{\widetilde{T}_i}\backslash^{\widetilde{T}_o}_{\square}\backslash_{p_{n+1}} \\ \pi'_n(x) & \triangleq & r_n\backslash^{\square}_{x_i}\backslash^{x_o}_{\square}\backslash^{\square}_{r_{n+1}}\backslash^{p_n}_{\square}\backslash_{p_{n+1}} \end{array}$$

Then we have

$$P_1 \triangleq P_1(a,G,C,T) \quad P_2 \triangleq P_2(a,C,G,T') \quad P_3 \triangleq P_3(F,T,T',C)$$

$$\begin{array}{lcl} P_4 & \triangleq & r_4\backslash^{\square}_{T'_i}\backslash^{T'_o}_{\square}\backslash^{\square}_{\overline{F}_i}\backslash^{\overline{F}_o}_{\square}\backslash^{\square}_{cxt}\backslash^{p_4}_{\square}\backslash^{\square}_{\widetilde{G}_i}\backslash^{\widetilde{G}_o}_{\square}\backslash_{\tau}.P_4 \\ & & + \\ & & r_4\backslash^{\square}_{\overline{T'}_i}\backslash^{\overline{T'}_o}_{\square}\backslash^{\square}_{cxt}\backslash^{p_4}_{\square}\backslash_{\tau}.P_4 \;+\; r_4\backslash^{\square}_{F_i}\backslash^{F_o}_{\square}\backslash^{\square}_{cxt}\backslash^{p_4}_{\square}\backslash_{\tau}.P_4 \end{array}$$

Encoding of the entities. As for reactions, also the encoding of the entities is given in a parametric way. Here we differentiate three types of encodings: (1) for the entities that are not provided by the context and can be produced by the reactions; (2) for the entities that can be provided by the context and can be produced by the reactions; (3) for the entities that are only provided by the context.

Here, the entities T and T' that can be produced by the reactions and that are not provided by the context:

$$P(T,\widetilde{T}) \triangleq \textstyle\sum_{h=0}^{1}(^{T_i}\backslash^{\square}_{T_o}\backslash\square)^h\ ^{\widetilde{T}_i}\backslash_{\widetilde{T}_o}.P(T,\widetilde{T}) \quad + \quad ^{T_i}\backslash_{T_o}.P(\overline{T},\widetilde{T})$$

Then, we have

$$\begin{array}{llllll} P_T & \triangleq & P(T,\widetilde{T}) & \overline{P}_T & \triangleq & P(\overline{T},\widetilde{T}) \quad P_{T'} \triangleq P(T',\widetilde{T'}) \\ \overline{P}_{T'} & \triangleq & P(\overline{T'},\widetilde{T'}) & & & \end{array}$$

The entities that can be produced by the reactions and that can be provided by the context are as follows:

$$\begin{array}{lcl} P(C,\widehat{C},\underline{C},\widetilde{C}) & \triangleq & \sum_{h=0}^{1}(^{C_i}\backslash^{\square}_{C_o}\backslash\square)^h\ ^{\widehat{C}_i}\backslash^{\square}_{\widehat{C}_o}\backslash\square(^{\widetilde{C}_i}\backslash^{\square}_{\widetilde{C}_o}\backslash\square)^h.P(C,\widehat{C},\underline{C},\widetilde{C}) \\ & & + \\ & & \sum_{h=0}^{1}(^{C_i}\backslash^{\square}_{C_o}\backslash\square)^h\ ^{\underline{C}_i}\backslash_{\underline{C}_o}.P(\overline{C},\widehat{C},\underline{C},\widetilde{C}) \\ & & + \\ & & \sum_{h=0}^{1}(^{C_i}\backslash^{\square}_{C_o}\backslash\square)^h\ ^{\underline{C}_i}\backslash_{\underline{C}_o}{}^{\square}\backslash^{\widehat{C}_i}_{\square}\backslash_{\widehat{C}_o}.P(C,\widehat{C},\underline{C},\widetilde{C}) \end{array}$$

Then, we have

$$P_C \triangleq P(C,\widehat{C},\underline{C},\widetilde{C}) \qquad P_G \triangleq P(G,\widehat{G},\underline{G},\widetilde{G})$$
$$\overline{P}_C \triangleq P(\overline{C},\widehat{C},\underline{C},\widetilde{C}) \qquad \overline{P}_G \triangleq P(\overline{G},\widehat{G},\underline{G},\widetilde{G})$$

The encoding of the entity F that can be provided by the context follows:

$$P_F \triangleq \sum_{h=0}^{1}({}^{F_i}\backslash^{\square}_{F_o}\backslash\square)^h\ {}^{\widehat{F}_i}\backslash_{\widehat{F}_o}.P_F \quad + \quad \sum_{h=0}^{1}({}^{F_i}\backslash^{\square}_{F_o}\backslash\square)^h.{}^{\underline{F}_i}\backslash_{\underline{F}_o}.\overline{P}_F$$
$$\overline{P}_F \triangleq \sum_{h=0}^{1}({}^{\overline{F}_i}\backslash^{\square}_{\overline{F}_o}\backslash\square)^h\ {}^{\widehat{F}_i}\backslash_{\widehat{F}_o}.P_F \quad + \quad \sum_{h=0}^{1}({}^{\overline{F}_i}\backslash^{\square}_{\overline{F}_o}\backslash\square)^h.{}^{\underline{F}_i}\backslash_{\underline{F}_o}.\overline{P}_F$$

Also in this example we account for a nondeterministic context that can (nondeterministically) provide any combination of the entities: C, G F:

$$Cxt \triangleq \sum_{\substack{C^\star \in \{C,\overline{C}\}\\ G^\star \in \{G,\overline{G}\}\\ F^\star \in \{F,\overline{F}\}}} {}^{cxt}\backslash^{\square}_{C^\star_i}\backslash^{C^\star_o}_{\square}\backslash^{\square}_{G^\star_i}\backslash^{G^\star_o}_{\square}\backslash^{\square}_{F^\star_i}\backslash^{F^\star_o}_{\square}\backslash_{p_1}.Cxt$$

Now, to show a possible composition of a transition label, we assume a system where only the entities a, T', and G are present:

$$Sys \triangleq I \mid P_a \mid \overline{P}_C \mid P_G \mid \overline{P}_F \mid \overline{P}_T \mid P_{T'} \mid P_1 \mid P_2 \mid P_3 \mid P_4 \mid Cxt$$

In the above configuration, reactions 1 and 4 can be applied, and also we assume that the context will provide the entity F, that will be available in the target configuration. Instead of showing the complete transition label, we give its flattened version obtained by applying the function $\mathit{flat}(\cdot)$:

$$r_1\ a\ G\ \overline{C}\ r_2\ G\ r_3\ T'\ r_4\ T'\ \overline{F}\ cxt\ \underline{C}\ \underline{G}\ \widehat{F}\ p_1\ \widetilde{T}\ p_2\ p_3\ p_4\ \widetilde{G}.$$

The original label can then be reconstructed just applying the function $\mathit{unflat}(\cdot)$ to the string above.

In [16] we have shown a more complex example, by modeling a RS of a regulatory network for *lac* operon, presented in [4].

6. Bio-simulation

The classical notion of bisimulation for process algebras equates two processes when one process can simulate all the instructions executed by the other one and viceversa. In its weak formulation, internal instructions, i.e. non visible by external observers, are abstracted away. There are many variants of the bisimulation for process algebras, for example the barbed bisimulation [18] only consider the execution of invisible actions, and then equates two processes when the expose the same prefixes; for the mobile ambients [10], a process algebra equipped with a reduction semantics, a notion of behavioural equivalence equates two processes when they expose the same ambients [19].

There are some previous works based on bisimulation applied to models for biological systems. Barbuti et al [20] define a classical setting for bisimulation

for two formalisms: the Calculus of Looping Sequences, which is a rewriting system, and the Brane Calculi, which is based on process calculi. Bisimulation is used to verify properties of the regulation of lactose degradation in Escherichia coli and the EGF signalling pathway. These calculi allow the authors to model membranes' behaviour. Cardelli et al [21] present two quantitative behavioral equivalences over species of a chemical reaction network with semantics based on ordinary differential equations. Bisimulation identifies a partition where each equivalence class represents the exact sum of the concentrations of the species belonging to that class. Bisimulation also relates species that have identical solutions at all time points when starting from the same initial conditions. Both the mentioned formalisms [20, 21] adopt a classical approach to bisimulation. Albeit the bisimulation is a powerful tool for verifying if the behaviour of two different software programs is indistinguishable, in the case of biological systems the classical bisimulation seems to be inappropriate, as it considers too many details. In fact, in a biological soup, a high number of interactions occur every seconds, and generally, biologists are only interested to analyse a small subset of them.

For this reason, we propose an alternative notion of bisimulation, that hereafter we call *bio-simulation*, that allows us to compare two biological systems by restricting the observation to only the limited events of interest. This allows one to tailor the equivalence to different applications and purposes.

The transition labels of our systems record detailed information about all the reactions that have been applied in one transition, about the elements that acted as reagents, as inhibitors or that have been produced, or that have been provided by the context. All these information are stored in the label because they are necessary to compose a transition in a modular way. Depending on the application, only a suitable abstraction over the label can be of interest.

In a way, at each step of the bisimulation game, we want to query our transition labels to get only the information we care about. To this goal, we introduce a simple language that allows us to formulate detailed and partial queries about what happened in a single transition.

Example 20. For instance we would like to express properties about each step of the bio-simulation of a system like the ones below:

1. Has the entity s_i been used by rule r_j as reagent?
2. Has the entity s_i blocked the application of rule r_j?
3. Has the entity s_i been produced by rule r_j?
4. Has the entity s_i been produced by some rule?
5. Has the entity s_i been provided by the context?
6. Has the rule r_j not been applied?

As detailed before, in the following we assume that: (i) the context can be non-deterministic, otherwise it makes little sense to rely on bisimulation to observe the branching structure of system dynamics; (ii) we are interested in observing the names of the entities involved in the transitions and also the rules that have been applied, thus we assume top level restrictions are absent and rely on solid transitions only (with leftmost and rightmost silent actions).

6.1. Assertion language

Next, we introduce an assertion language that operates on simple strings and that combines regular expression operators with conjunction and disjunction. We let $\mathit{syms} = \mathit{decs} \cup \mathit{reacts} \cup \mathit{prods} \cup \{\mathit{cxt}\}$ be the set of symbolic names used in our assertion language.

Definition 21 (Assertion Language). Assertions are built from:

$$\begin{array}{lcl} \zeta & ::= & \alpha \mid ? \mid [N] \\ \mathsf{F} & ::= & \epsilon \mid \zeta \mid \mathsf{F} :: \mathsf{F} \mid \mathsf{F}^+ \mid \mathsf{F}^* \mid \mathsf{F} \vee \mathsf{F} \mid \mathsf{F} \wedge \mathsf{F} \end{array}$$

where $\alpha \in \mathit{syms}$ and $N \subseteq \mathit{syms}$.

Roughly, a *singleton* assertion ζ denotes either a string composed by a single symbol α (one of the symbols in the set *syms* for denoting a particular entity, rule, production or context), or the wildcard ? that stands for any symbol, or the pattern $[N]$ that stands for any of the strings composed by a single symbol in the set N. Clearly ? is just a shorthand for $[\mathit{syms}]$. We write $\mathbf{0}$ for $[\emptyset]$ and $[s_1, ..., s_n]$ instead of $[\{s_1, ..., s_n\}]$.

An *assertion* F is either the empty string ϵ, a singleton assertion ζ, the concatenation of two assertions $\mathsf{F}_1 :: \mathsf{F}_2$, the replication of F for 1 or more times F^+, the replication of F for 0 or more times F^*, the disjunction of two assertions $\mathsf{F}_1 \vee \mathsf{F}_2$ or their conjunction $\mathsf{F}_1 \wedge \mathsf{F}_2$. We write $\star$ as an abbreviation of $?^*$.

An assertion denotes a set of strings over the alphabet *syms* as expected.

Definition 22 (Semantics of Assertions). We define $[\![\mathsf{F}]\!] \subseteq \wp(\mathit{syms}^*)$ by induction on the structure of F:

$$\begin{array}{rcl@{\qquad}rcl} [\![\epsilon]\!] & \triangleq & \{\epsilon\} & [\![\mathsf{F}_1 :: \mathsf{F}_2]\!] & \triangleq & \{\omega_1 :: \omega_2 \mid \omega_1 \in [\![\mathsf{F}_1]\!] \wedge \omega_2 \in [\![\mathsf{F}_2]\!]\} \\ [\![\alpha]\!] & \triangleq & \{\alpha\} & [\![\mathsf{F}^+]\!] & \triangleq & [\![\mathsf{F}]\!]^+ \\ [\![?]\!] & \triangleq & \mathit{syms} & [\![\mathsf{F}^*]\!] & \triangleq & [\![\mathsf{F}]\!]^* \\ [\![[N]]\!] & \triangleq & N & [\![\mathsf{F}_1 \vee \mathsf{F}_2]\!] & \triangleq & [\![\mathsf{F}_1]\!] \cup [\![\mathsf{F}_2]\!] \\ & & & [\![\mathsf{F}_1 \wedge \mathsf{F}_2]\!] & \triangleq & [\![\mathsf{F}_1]\!] \cap [\![\mathsf{F}_2]\!] \end{array}$$

Definition 23 (Satisfaction as Membership). Let υ be a transition label, and F be an assertion. We write $\upsilon \models \mathsf{F}$ (read as the transition label υ satisfies the assertion F) if $\mathit{flat}(\upsilon) \in [\![\mathsf{F}]\!]$, otherwise we write $\upsilon \not\models \mathsf{F}$ (or also $\upsilon \models \neg\mathsf{F}$) and say that F does not hold at υ.

Given two transition labels v, w we write $v \equiv_{\mathsf{F}} w$ if $v \models \mathsf{F} \Leftrightarrow w \models \mathsf{F}$, i.e. if both v, w satisfy F or they do not.

Example 24. The formulas corresponding to the sample queries listed in Example 20 are as follows:

1. $\star :: r_j :: [s_1, ..., s_n]^* :: s_i :: [s_1, ..., s_n]^* :: [\overline{s}_1, ..., \overline{s}_n]^+ :: \star$
2. $\star :: r_j :: [s_i, \overline{s}_i] :: r_{j+1} :: \star$
3. $\star :: p_j :: [\mathit{ents}]^* :: \widetilde{s}_i :: \star$

4. $\star :: \widetilde{s}_i :: \star$
5. $\star :: \widehat{s}_i :: \star$
6. $\star :: r_j ::? :: r_{j+1} :: \star$

where in 1, 2, 6 we exploit the fact that in a reaction (R, I, P) the sets R of reactants and I of inhibitors are non empty, so that if there is only one symbol between the occurrence of r_j and r_{j+1} it means the reaction r_j has not been applied. Viceversa, if the reaction r_j has been applied the occurrence of r_j must be followed by at least one of the symbols in $\{s_1, ..., s_n\}$ and then by at least one of the symbols in $\{\overline{s}_1, ..., \overline{s}_n\}$.

6.2. Bio-similarity and bio-logical equivalence

The notion of bio-simulation builds on the above language of assertions to parameterize the induced equivalence on the property of interest. Please recall that we have defined the behaviour of the contest in a non deterministic way, thus at each step, different possible sets of entities can be provided to the system and different sets of reaction can be enabled/disabled. Bio-simulation can thus be used to compare the behaviour of different systems that share some of the reactions or entities or also to compare the behaviour of the same set of reaction rules when different contexts are provided.

Definition 25 (Bio-similarity $\sim_{\mathsf{F}}$). Given an assertion F, a *bio-simulation* $\mathbf{R}_{\mathsf{F}}$ that respects F is a binary relation over cCNA processes such that, whenever $P \ \mathbf{R}_{\mathsf{F}} \ Q$ then:

- for any v, P' such that $P \stackrel{v}{\Rightarrow} P'$ then there exist w, Q' such that $Q \stackrel{w}{\Rightarrow} Q'$ with $v \equiv_{\mathsf{F}} w$ and $P' \ \mathbf{R}_{\mathsf{F}} \ Q'$.
- for any w, Q' such that $Q \stackrel{w}{\Rightarrow} Q'$ then there exist v, P' such that $P \stackrel{v}{\Rightarrow} P'$ with $v \equiv_{\mathsf{F}} w$ and $P' \ \mathbf{R}_{\mathsf{F}} \ Q'$.

We let $\sim_{\mathsf{F}}$ denote the largest bio-simulation and we say that P is *bio-similar* to Q, with respect to F, if $P \sim_{\mathsf{F}} Q$.

Remark 26. Please remember that the notation $P \stackrel{v}{\Rightarrow} P'$ refers to ordinary transitions $P \stackrel{v}{\rightarrow} P'$ where v is a complete chain (solid and with τ actions at the extremes). The double arrow notation should not be confused with the notation for weak transitions commonly found in the literature on process algebras.

Remark 27. An alternative way to look at a bio-simulation that respects F is to define it as an ordinary bisimulation over the transition system labelled over $\{\mathsf{F}, \neg\mathsf{F}\}$ obtained by transforming each transition $P \stackrel{v}{\Rightarrow} P'$ such that $v \models \mathsf{F}$ into $P \stackrel{\mathsf{F}}{\rightarrow} P'$ and each transition $P \stackrel{v}{\Rightarrow} P'$ such that $v \not\models \mathsf{F}$ into $P \stackrel{\neg\mathsf{F}}{\longrightarrow} P'$.

It can be easily shown that the identity relation is a bio-simulation and that bio-simulations are closed under (relational) inverse, composition and union and that, as a consequence, bio-similarity is an equivalence relation.

Now, we introduce a slightly modified version of the Hennessy Milner Logic [13], called bioHML; due to the reasons we explained above, we do not want to look at the complete transition labels, thus we rely on our simple assertion language to make it parametric to the assertion F of interest:

Definition 28 (BioHML). Let F be an assertion, then the set of bioHML formulas G that respects F are built by the following syntax:

$$\begin{array}{rcl} \chi & ::= & \mathsf{F} \mid \neg\mathsf{F} \\ \mathsf{G},\mathsf{H} & ::= & \mathtt{t} \mid \mathtt{f} \mid \mathsf{G}\wedge\mathsf{G} \mid \mathsf{G}\vee\mathsf{G} \mid \langle\chi\rangle\mathsf{G} \mid [\chi]\mathsf{G} \end{array}$$

Remark 29. An alternative way to look at bioHML formulas is as ordinary HML formulas over the set of labels $\{\mathsf{F}, \neg\mathsf{F}\}$.

As usual, the semantics of a bioHML formula is the set of processes that satisfy it.

Definition 30 (Semantics of BioHML). We define $[\![\mathsf{G}]\!] \subseteq \mathcal{P}$ by induction on the structure of G:

$$\begin{array}{lcllcl} [\![\mathtt{t}]\!] & \triangleq & \mathcal{P} & [\![\mathsf{G}\wedge\mathsf{H}]\!] & \triangleq & [\![\mathsf{G}]\!]\cap[\![\mathsf{H}]\!] \\ [\![\mathtt{f}]\!] & \triangleq & \emptyset & [\![\mathsf{G}\vee\mathsf{H}]\!] & \triangleq & [\![\mathsf{G}]\!]\cup[\![\mathsf{H}]\!] \end{array}$$

$$\begin{array}{lcl} [\![\langle\chi\rangle\mathsf{G}]\!] & \triangleq & \{P\in\mathcal{P} : \exists v, P'.\ P \stackrel{v}{\Rightarrow} P' \text{ with } v\models\chi \text{ and } P'\in[\![\mathsf{G}]\!]\} \\ [\![[\chi]\mathsf{G}]\!] & \triangleq & \{P\in\mathcal{P} : \forall v, P'.\ P \stackrel{v}{\Rightarrow} P' \text{ implies } v\models\chi \text{ and } P'\in[\![\mathsf{G}]\!]\} \end{array}$$

We write $P \models \mathsf{G}$ (P satisfies G) if and only if $P \in [\![\mathsf{G}]\!]$.

Negation is not included in the syntax, but the converse $\overline{\mathsf{G}}$ of a bioHML formula G can be easily defined inductively in the same way as for HML logic.

Definition 31 (Converse). Given a bioHML formula G we define its converse $\overline{\mathsf{G}}$ as follows:

$$\begin{array}{lcllcllcl} \overline{\mathtt{t}} & \triangleq & \mathtt{f} & \overline{\mathsf{G}\wedge\mathsf{H}} & \triangleq & \overline{\mathsf{G}}\vee\overline{\mathsf{H}} & \overline{\langle\chi\rangle\mathsf{G}} & \triangleq & [\chi]\overline{\mathsf{G}} \\ \overline{\mathtt{f}} & \triangleq & \mathtt{t} & \overline{\mathsf{G}\vee\mathsf{H}} & \triangleq & \overline{\mathsf{G}}\wedge\overline{\mathsf{H}} & \overline{[\chi]\mathsf{G}} & \triangleq & \langle\chi\rangle\overline{\mathsf{G}} \end{array}$$

We observe that, as expected, for any bioHML formula G and process P we have $\overline{\overline{\mathsf{G}}} = \mathsf{G}$ and $P \models \overline{\mathsf{G}}$ iff $P \not\models \mathsf{G}$.

We let $\mathcal{L}_\mathsf{F}$ be the set of all bioHML formulas that respects F and we say that two processes P, Q are bio-logically equivalent w.r.t. F, written $P \equiv_{\mathcal{L}_\mathsf{F}} Q$, when P and Q satisfy the exactly the same bioHML formulas in $\mathcal{L}_\mathsf{F}$, i.e. when for any $\mathsf{G}\in\mathcal{L}_\mathsf{F}$ we have $P\models\mathsf{G} \Leftrightarrow Q\models\mathsf{G}$.

Finally, we extend the classical result establishing the correspondence between the logical equivalence induced by HML with bisimilarity for proving that bio-similarity coincides with bio-logical equivalence.

Theorem 32 (Correspondence). $\sim_\mathsf{F} = \equiv_{\mathcal{L}_\mathsf{F}}$

6.3. Bio-simulation at work

We will show how bio-simulation works. For the sake of space, we consider a very simple example with only two reactions. Reaction P_1 requires G to produce C; reaction P_2 requires C to produce G; both reactions have H as inhibitor. Now, we set two systems defined by the same two reactions, with the two different initial configuration, and with two different context definitions. The two reactions work as follows (where we omit to specify the cases where H is present, as they will never happen):

$$\begin{array}{rcl} P_1 & \triangleq & \tau\backslash^{\square}_{G_i}\backslash^{G_o}_{\square}\backslash^{\square}_{\overline{H}_i}\backslash^{\overline{H}_o}_{\square}\backslash^{\square}_{r_2}\backslash^{p_1}_{\square}\backslash^{\square}_{\widetilde{C}_i}\backslash^{\widetilde{C}_o}_{\square}\backslash_{p_2}.P_1 \quad + \quad \tau\backslash^{\square}_{\overline{G}_i}\backslash^{\overline{G}_o}_{\square}\backslash^{\square}_{r_2}\backslash^{p_1}_{\square}\backslash_{p_2}.P_1 \\ P_2 & \triangleq & r_2\backslash^{\square}_{C_i}\backslash^{C_o}_{\square}\backslash^{\square}_{\overline{H}_i}\backslash^{\overline{H}_o}_{\square}\backslash^{\square}_{cxt}\backslash^{p_2}_{\square}\backslash^{\square}_{\widetilde{G}_i}\backslash^{\widetilde{G}_o}_{\square}\backslash_{\tau}.P_2 \quad + \quad r_2\backslash^{\square}_{\overline{C}_i}\backslash^{\overline{C}_o}_{\square}\backslash^{\square}_{cxt}\backslash^{p_2}_{\square}\backslash_{\tau}.P_2 \end{array}$$

The two contexts follow :

$$Cxt_1 \triangleq {}^{cxt}\backslash^{\square}_{\widehat{C}_i}\backslash^{\widehat{C}_o}_{\square}\backslash^{\square}_{\underline{G}_i}\backslash^{\underline{G}_o}_{\square}\backslash^{\square}_{\underline{H}_i}\backslash^{\underline{H}_o}_{\square}\backslash_{p_1}.Cxt_1 + {}^{cxt}\backslash^{\square}_{\underline{C}_i}\backslash^{\underline{C}_o}_{\square}\backslash^{\square}_{\underline{G}_i}\backslash^{\underline{G}_o}_{\square}\backslash^{\square}_{\underline{H}_i}\backslash^{\underline{H}_o}_{\square}\backslash_{p_1}.Cxt_1$$

$$Cxt_2 \triangleq {}^{cxt}\backslash^{\square}_{\widehat{G}_i}\backslash^{\widehat{G}_o}_{\square}\backslash^{\square}_{\underline{C}_i}\backslash^{\underline{C}_o}_{\square}\backslash^{\square}_{\underline{H}_i}\backslash^{\underline{H}_o}_{\square}\backslash_{p_1}.Cxt_2 + {}^{cxt}\backslash^{\square}_{\underline{G}_i}\backslash^{\underline{G}_o}_{\square}\backslash^{\square}_{\underline{C}_i}\backslash^{\underline{C}_o}_{\square}\backslash^{\square}_{\underline{H}_i}\backslash^{\underline{H}_o}_{\square}\backslash_{p_1}.Cxt_2$$

The definition of the processes encoding G and C is similar, and it is given in a parametric way:

$$P(G) \triangleq {}^{G_i}\backslash_{G_o}.\overline{P}(G) \qquad \overline{P}(G) \triangleq {}^{\overline{G}_i}\backslash_{\overline{G}_o}{}^{\square}\backslash_{\square}{}^{\widetilde{G}_i}\backslash_{\widetilde{G}_o}P(G)$$

and we have $P_G \triangleq P(G)$, $P_C \triangleq P(C)$, then $\overline{P}_H \triangleq {}^{\overline{H}_i}\backslash_{\overline{H}_o}.\overline{P}_H + {}^{\overline{H}_i}\backslash^{\square}_{\overline{H}_o}\backslash^{\overline{H}_i}_{\square}\backslash_{\overline{H}_i}.\overline{P}_H$, as H is neither produced nor provided by the context. Then, the initial configuration of system Sys_1 includes C and not G and the context can only provide G, the initial configuration of system Sys_2 includes G and not C and the context can only provide C:

$$\begin{array}{rcl} Sys_1 & \triangleq & I \mid P_1 \mid P_2 \mid P_G \mid \overline{P}_C \mid \overline{P}_H \mid Cxt_1 \\ Sys_2 & \triangleq & I \mid P_1 \mid P_2 \mid \overline{P}_G \mid P_C \mid \overline{P}_H \mid Cxt_2 \end{array}$$

Fig. 4 shows the labelled transition system of Sys_1, with initial state only containing G, and the transition system of Sys_2, with initial state only containing C, limited to the complete and solid labels. As before we show the output of the *flat*$(\cdot)$ function applied to the transition labels:

$$\begin{array}{rcl} \ell_1 & \triangleq & r_1\ G\ \overline{H}\ r_2\ \overline{C}\ cxt\ \widehat{C}\ \underline{G}\ \underline{H}\ p_1\ \widetilde{C}\ p_2 \\ \ell_2 & \triangleq & r_1\ \overline{G}\ r_2\ C\ \overline{H}\ cxt\ \widehat{C}\ \underline{G}\ \underline{H}\ p_1\ p_2\ \widetilde{G} \end{array} \qquad \begin{array}{rcl} \ell'_1 & \triangleq & r_1\ G\ \overline{H}\ r_2\ \overline{C}\ cxt\ \underline{C}\ \underline{G}\ \underline{H}\ p_1\ \widetilde{C}\ p_2 \\ \ell'_2 & \triangleq & r_1\ \overline{G}\ r_2\ C\ \overline{H}\ cxt\ \underline{C}\ \underline{G}\ \underline{H}\ p_1\ p_2\ \widetilde{G} \end{array}$$

$$\begin{array}{rcl} \ell_3 & \triangleq & r_1\ G\ \overline{H}\ r_2\ C\ \overline{H}\ cxt\ \widehat{C}\ \underline{G}\ \overline{H}\ p_1\ \widetilde{C}\ p_2\ \widetilde{G} \\ \ell'_3 & \triangleq & r_1\ G\ \overline{H}\ r_2\ C\ \overline{H}\ cxt\ \underline{C}\ \underline{G}\ \overline{H}\ p_1\ \widetilde{C}\ p_2\ \widetilde{G} \end{array}$$

The labels $\flat_i$ $\flat'_i$, with $i \in \{1,2,3\}$ can be obtained by labels ℓ_i, ℓ'_i by substituting G with C and viceversa.

Now, it easy to check that Sys_1 and Sys_2 are bio-similar w.r.t the property F

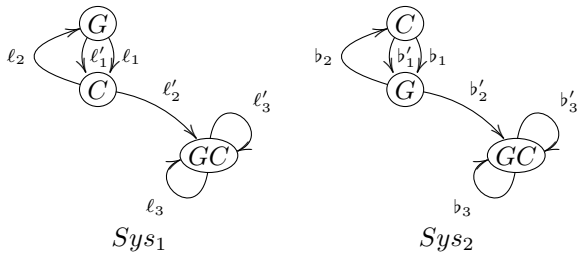

Figure 4: The two labelled transition systems of Sys_1 and Sys_2.

saying that G and C are simultaneously produced, formally: $Sys_1 \sim_{\mathsf{F}} Sys_2$ with $\mathsf{F} = \star :: \widetilde{G} :: \star \wedge \star :: \widetilde{C} :: \star$.

On the contrary, $Sys_1 \not\sim_{\mathsf{F}'} Sys_2$ with $\mathsf{F}' = \star :: \widetilde{C} :: \star$, because it happens that both transition labels ℓ_1 and ℓ'_1, in Sys_1, record the production of C, whereas transition labels $\flat_1$ and $\flat'_1$ do not. In fact, the bioHML formula $\mathsf{G} \triangleq \langle \mathsf{F}' \rangle \mathtt{t}$ can be used to distinguish Sys_1 from Sys_2, as $Sys_1 \models \mathsf{G}$ and $Sys_2 \not\models \mathsf{G}$.

7. Enhanced Reaction Systems

Our encoding increases the expressivity of RS concerning: the possibility of alternative behaviour of mutated entities, and the communication between two different Reaction Systems. It is important to note that, when the context is deterministic, our encoding guarantees that from each state, in the cCNA transition system, only one state is reachable, as the dynamics is totally deterministic.

7.1. Mutating entities

In RS, when an entity is present, it can potentially be involved in each reactions where it is required. With a few more lines of code, in cCNA it is possible to describe the behaviour of a mutation of an entity, in a way that the mutated version of the entity can take part to only a subset of the rules requiring the *normal version* of the entity. For example, let us assume that entity $s1$ is consumed by reactions $a1$ and $a2$. Reaction $a1$ produces also $s1$ if $s2$ is present, otherwise $a1$ produces a mutated version of $s1$, say $s1'$. When $s1'$ is produced, reaction $a2$ behaves in the same way as if $s1$ would be absent, whereas $a2$ recognises the presence of $s1'$ and behaves in the same way as if $s1$ would be present. Technically, in both cases it is enough to add one more nondeterministic choice in the code of P_{a1} and P_{a2}.

7.2. Communicating Reaction Systems

We sketch how it is possible to program two RSs encodings, in a way that the entities that usually come from the context of one RS will be provided instead from the other RS.

$rs1$	$rs2$
$a_1 = (s, \ , x)$	$a_2 = (y, \ , s)$

Table 1: The two Reaction Systems $rs1$ and $rs2$.

Example 33. Let $rs1$ and $rs2$ be two RSs, defined by the rules in Table 1. Now, we set our example such that the two contexts, for $rs1$ and $rs2$, do not provide any entities. We also assume that entity s in $rs1$ is provided by $rs2$, as $rs2$ produces a quantity of s that is enough for $rs1$ and $rs2$. For technical reasons, we can not use the same name for s in both the two RSs, then we use the name ss in $rs2$. We need to modify our translation technique to suite this new setting. As we do not model contexts, we introduce *dummy* channel names dx and dss to model the absence of entities. Also, thanks to the simplicity of the example, we can leave out the use of the p_i channels. This streamlining does not affect the programming technique we propose to make two RSs communicate. First, we translate the reaction in $rs1$:

$$[\![a_1]\!] \triangleq P_{a_1} \quad \triangleq \quad \tau\backslash^{\square}_{s_i}\backslash^{s_o}_{\square}\backslash^{\square}_{\widetilde{x}_i}\backslash^{\widetilde{x}_o}_{\square}\backslash_{a_2}.P_{a_1} + \tau\backslash^{\square}_{\overline{s}_i}\backslash^{\overline{s}_o}_{\square}\backslash^{\square}_{dx_i}\backslash^{dx_o}_{\square}\backslash_{a_2}.P_{a_1}$$

Please note, that prefixes of process P_{a_1} end with the channel name a_2, as the link chain is now connected with the reaction of $rs2$. The translation for the entities follows.

$$\begin{array}{rcccl}
[\![s]\!] & \triangleq & P_s & \triangleq & {}^{s_i}\backslash^{\square}_{s_o}\backslash^{\widehat{s}_i}_{\square}\backslash_{\widehat{s}_o}.P_s + {}^{s_i}\backslash_{s_o}.\overline{P_s} \\
 & & \overline{P_s} & \triangleq & {}^{\overline{s}_i}\backslash^{\square}_{\overline{s}_o}\backslash^{\widehat{s}_i}_{\square}\backslash_{\widehat{s}_o}.P_s + {}^{\overline{s}_i}\backslash_{\overline{s}_i}.\overline{P_s} \\
[\![x]\!] & \triangleq & P_x & \triangleq & {}^{\widetilde{x}_i}\backslash_{\widetilde{x}_o}.P_x + {}^{dx_i}\backslash_{dx_o}.\overline{P_x} \\
 & & \overline{P_x} & \triangleq & {}^{\widetilde{x}_i}\backslash_{\widetilde{x}_o}.P_x + {}^{dx_i}\backslash_{dx_o}.\overline{P_x}
\end{array}$$

The translation for $rs2$ follows.

$$[\![a_2]\!] \triangleq P_{a_2} \quad \triangleq \quad a_2\backslash^{\square}_{y_i}\backslash^{y_o}_{\square}\backslash^{\square}_{\widetilde{ss}_i}\backslash^{\widetilde{ss}_o}_{\square}\backslash_{\tau}.P_{a_2} + {}^{a_2}\backslash^{\square}_{\overline{y}_i}\backslash^{\overline{y}_o}_{\square}\backslash^{\square}_{dss_i}\backslash^{dss_o}_{\square}\backslash_{\tau}.P_{a_2}$$

In the translation of the entities in $rs2$, we introduce the mechanism that allows the entity s (ss in $rs2$) to be provided in $rs1$. Every time ss is produced in $rs2$, a virtual link is created to synchronise with $rs1$ on link ${}^{\widehat{s}_i}\backslash_{\widehat{s}_o}$:

$$\begin{array}{rcccl}
[\![ss]\!] & \triangleq & P_{ss} & \triangleq & {}^{\widetilde{ss}_i}\backslash^{\square}_{\widehat{s}_i}\backslash^{\widehat{s}_o}_{\square}\backslash_{\widetilde{ss}_o}.P_{ss} + {}^{dss_i}\backslash_{dss_o}.\overline{P_{ss}} \\
 & & \overline{P_{ss}} & \triangleq & {}^{\widetilde{ss}_i}\backslash^{\square}_{\widehat{s}_i}\backslash^{\widehat{s}_o}_{\square}\backslash_{\widetilde{ss}_o}.P_{ss} + {}^{dss_i}\backslash_{dss_o}.\overline{P_{ss}} \\
[\![y]\!] & \triangleq & P_y & \triangleq & {}^{y_i}\backslash_{y_o}.\overline{P_y} \\
 & & \overline{P_y} & \triangleq & {}^{\overline{y}_i}\backslash_{\overline{y}_o}.\overline{P_y}
\end{array}$$

We now assume that the initial system is $S \triangleq (\nu\, names)(P_{a_1}|P_{a_2}|P_s|P_y|\overline{P_x}|\overline{P_{ss}})$, i.e. only entities s and y are present. Now, the only possible transition has the following label (that we report without restriction):

$$\tau\backslash^{s_i}_{s_i}\backslash^{s_o}_{s_o}\backslash^{\widetilde{x}_i}_{\widetilde{x}_i}\backslash^{\widetilde{x}_o}_{\widetilde{x}_o}\backslash^{a_2}_{a_2}\backslash^{y_i}_{y_i}\backslash^{y_o}_{y_o}\backslash^{\widetilde{ss}_i}_{\widetilde{ss}_i}\backslash^{\widehat{s}_i}_{\widehat{s}_i}\backslash^{\widehat{s}_o}_{\widehat{s}_o}\backslash^{\widetilde{ss}_o}_{\widetilde{ss}_o}\backslash_{\tau},$$

where the black links belong to the prefixes of P_{a_1}, and P_{a_2}, the blue links belong to P_s, the gray links belong to P_y, and $\overline{P_x}$ and the red links belong to $\overline{P_{ss}}$. After the execution, the entity s is still present in $rs1$ as it has been provided by $rs2$.

As we have briefly sketched, our model of two *communicating Reaction Systems* can enable the study of the behaviour of one RS in relation to another one. Thus, the products of the reactions of one RS can become the input for another one. This could allow for a modular approach to modeling complex systems, by composing different Reaction Systems.

8. Conclusion

In this paper we have introduced cCNA, a variant of the `link`-calculus where prefixes are link chains and no more single links, as it was briefly described in the future work section in [9]. This variant allowed us to define a faithful embedding of Reaction Systems, an emerging formalism to model computationally biochemical systems. This translation shows several benefits. For instance, in our paper the context of a RS can work non deterministically and it is recursively defined. Also, entity mutations can be expressed easily and different reaction systems can communicate between them.

We have then modified the classical notion (in concurrency) of bisimulation making it more suitable for proving properties of Reaction Systems in our framework. We have defined a new assertion language, which allows us to specify the properties to be verified, and have shown that it extends Hennessy-Milner logic to our framework. We believe that our work can make possible to investigate how to integrate our methodology with other formal techniques to prove properties of the modeled systems [22, 23, 24].

We are confident that our embedding can contribute to extend the applications of Reaction Systems to diverse fields of computer science, and life sciences.

As future work, we plan to implement a prototype of our framework, with an automatic translation from RSs to `link`-calculus. We can also exploit the implementation of the symbolic semantics of the `link`-calculus [25] that can be found in [26].

We also plan to build on the work in this paper to extend the framework of RSs towards a model where different RSs are allowed to communicate.

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Appendix A. Omitted Proofs

In this section we report the proofs for the results in Section 4 and in Section 6.

Lemma 12. *Let $\mathcal{A} = (S, A)$ be a RS and let $\pi = (\gamma, \delta)$ be an extended interactive process in $\mathcal{A}$. Let $P = [\![\mathcal{A}, \gamma]\!]$ its cCNA translation. If exists P' such that $P \xrightarrow{(\nu\, names)\upsilon} P'$ is a transition of P, then*

1. *for each reaction $aj \in A$, the corresponding channels r_j and p_j appear in υ; for each entity $s \in S$, the corresponding channel s (suitably decorated) appear in υ; the channel cxt appears in υ;*
2. *for each reaction $aj \in A$ and each entity $s \in S$, each virtual link offered by processes P_a and Cxt is overlapped by exactly one solid link offered by processes representing entities.*

Proof. We prove the two items separately:

1. by Definition 11, all the names that appear in the prefixes of any subprocess are in the set *names* and thus restricted. They include all the reaction names r_j, all the production names p_j, all the entity names s_i, s_o, in all their decorated versions, and the name cxt. Therefore the chain υ must start and end with a τ action and cannot contain virtual links. The only prefix that starts with τ is the one of the recursive init process I (prefix ${}^{\tau}\backslash_{r_1}$) and the only prefixes that end with τ are those associated to reaction a_u (as we have assumed that $p_{u+1} = \tau$, where u is the number of reactions). Then each prefix that starts with r_j involves p_j and r_{j+1}. Thus all the prefixes associated with reactions must be concatenated and also the prefix associated with the context (remember that $r_{u+1} = cxt$), forming the backbone of the label. Since the context processes is involved, then all entities processes are also involved. Then, all the processes I, P_a, P_s (or $\overline{P_s}$), and Cxt must participate to each transition.
2. for each reaction $aj \in A$, the cCNA code of P_{aj} leaves one virtual link between two solid links of the types ${}^{r_j}\backslash_{\dots}\backslash_{s_i}^{\square}\backslash_{\square}^{s_o}\backslash_{\dots}\backslash_{r_{j+1}}^{\dots}{}^{p_j}\backslash_{\dots}\backslash_{p_{j+1}}$. Then, it derives that the process P_s, encoding the behaviour of entity s, can participate by filling the virtual link in the above transition by only offering one solid link of the form ${}^{s_i}\backslash_{s_o}$. In fact, there is no other way to generate a solid chain from s_i to s_o. The same reasoning holds for the processes Cxt and for all the decorated versions of s_i, s_o.

Proposition 16 (Correctness 1). *Let $P = [\![\mathcal{A}, \gamma]\!]$ with*

$$P = (\nu\, names)\left(I \mid \prod_{a \in A} P_a \mid Cxt^1 \mid \prod_{s \in C_0} P_s \mid \prod_{s \notin C_0} \overline{P}_s\right).$$

If there exists P' such that $P \xrightarrow{\upsilon} P'$, it holds that:

1. $v = {}^{\tau}\backslash_{\tau} \ldots {}^{\tau}\backslash_{\tau}$, *and*
2. $P' = (\nu\, names)\left(I \mid \prod_{a\in A} P_a \mid Cxt^2 \mid \prod_{s\in C_1\cup D_1} P_s | \prod_{s\notin C_1\cup D_1} \overline{P}_s\right)$.

Moreover, given $\pi^1 = (\gamma^1, \delta^1)$, *we have* $P' = [\![\mathcal{A}, \gamma^1]\!]$.

PROOF. First, we note that all the channels in the system are restricted, see Def. 11, then it holds that the transition labels are of the form $v = {}^{\tau}\backslash_{\tau} \ldots {}^{\tau}\backslash_{\tau}$. Now, by Definition 11 and by Lemma 12.1, all the channels r_j, p_j, with $j \in [1, \ldots, u]$, and cxt_h, with $h \in [1, \ldots, w]$, and all the annotated versions of s_i, s_o are restricted. Also, processes Cxt always requires the interaction with P_s on either on channels $\widehat{s}_i$, $\widehat{s}_o$ or on channels $\underline{s}_i$, $\underline{s}_o$. It derives that all the processes: P_a (coding the behaviour of reaction $a \in A$), P_s (coding the behaviour of entity $s \in S$), and Cxt (coding the behaviour of the context regarding all the entities) have been involved in the transition.

For any process P_{aj} encoding a reaction aj we have the following cases:

(a) if aj is applicable and it produces the entity s, the process P_{aj} provides a code of this type:

$$P_{aj} \triangleq {}^{r_j}\backslash_{\ldots}\backslash^{\square}_{r_{j+1}}\backslash^{p_j}_{\square}\backslash_{\ldots}\backslash^{\square}_{\widetilde{s}_i}\backslash^{\ldots\widetilde{s}_o}_{\square}\backslash_{\ldots}\backslash_{p_{j+1}}.P_{aj};$$

(b) if aj is applicable and it consumes the entity s, the process P_{aj} provides a code of this type:

$$P_{aj} \triangleq {}^{r_j}\backslash_{\ldots}\backslash^{\square}_{s_i}\backslash_{\square}{}^{s_o}\backslash_{\ldots}\backslash^{\square}_{r_{j+1}}\backslash^{p_j}_{\square}\backslash_{\ldots}\backslash_{p_{j+1}}.P_{aj};$$

(c) if aj is applicable and it requires the absence of the entity s, the process P_{aj} provides a code of this type:

$$P_{aj} \triangleq {}^{r_j}\backslash_{\ldots}\backslash^{\square}_{\overline{s}_i}\backslash_{\square}{}^{\overline{s}_o}\backslash_{\ldots}\backslash^{\square}_{r_{j+1}}\backslash^{p_j}_{\square}\backslash_{\ldots}\backslash_{p_{j+1}}.P_{aj};$$

(d) if aj is not applicable, the process P_{aj} executes a code capturing either the absence of one of its reactants (case 1), or the presence of one of its inhibitors (case 2):

1. $P_{aj} \triangleq {}^{r_j}\backslash_{\ldots}\backslash^{\square}_{\overline{s}_i}\backslash_{\square}{}^{\overline{s}_o}\backslash_{\ldots}\backslash^{\square}_{r_{j+1}}\backslash^{p_j}_{\square}\backslash_{\ldots}\backslash_{p_{j+1}}.P_{aj}$;
2. $P_{aj} \triangleq {}^{r_j}\backslash_{\ldots}\backslash^{\square}_{s_i}\backslash_{\square}{}^{s_o}\backslash_{\ldots}\backslash^{\square}_{r_{j+1}}\backslash^{p_j}_{\square}\backslash_{\ldots}\backslash_{p_{j+1}}.P_{aj}$.

Now, we consider the structure of the process Cxt^1. By Definition 11, Cxt^1 is the unique process encoding the behaviour of the context regulating the supply of any entity.

(e) The code of the process Cxt^1 that provides s and not e has the following structure:

$$Cxt^1 \triangleq {}^{cxt}\backslash_{\ldots}\backslash^{\square}_{\widehat{s}_i}\backslash_{\square}{}^{\widehat{s}_o}\backslash_{\ldots}\backslash^{\square}_{\underline{e}_i}\backslash_{\square}{}^{\underline{e}_o}\backslash_{\ldots}\backslash_{p_1}.Cxt^2.$$

The code executed by P_s has the following structure:

(f) $P_s \triangleq \sum_{h,k\geq 0}({}^{s_i}\backslash^{\square}_{s_o}\backslash\square)^h\, {}^{\widehat{s}_i}\backslash^{\square}_{\widehat{s}_o}\backslash\square({}^{\widetilde{s}_i}\backslash^{\square}_{\widetilde{s}_o}\backslash\square)^k.P_s$, if $s \in C_{i+1}$;

(g) $P_s \triangleq \sum_{h\geq 0,k\geq 1}({}^{s_i}\backslash^{\square}_{s_o}\backslash\square)^h\, {}^{\underline{s}_i}\backslash^{\square}_{\underline{s}_o}\backslash\square({}^{\widetilde{s}_i}\backslash^{\square}_{\widetilde{s}_o}\backslash\square)^k.P_s$, if $s \notin C_{i+1}$

(h) $P_s \triangleq \sum_{h\geq 0}({}^{s_i}\backslash^{\square}_{s_o}\backslash\square)^h {}^{\underline{s}_i}\backslash_{\underline{s}_o}.\overline{P}_s$, if $s \notin C_{i+1}$

where, by Lemma 12.2, h is the number of reactions requiring the presence of s plus possibly some reactions not requiring s; and k is the number of reactions producing s.

Similarly, the code executed by $\overline{P}_s$ has the following structure:

(f') $\overline{P}_s \triangleq \sum_{h,k\geq 0}({}^{\overline{s}_i}\backslash^{\square}_{\overline{s}_o}\backslash\square)^h\, {}^{\widehat{s}_i}\backslash^{\square}_{\widehat{s}_o}\backslash\square({}^{\widetilde{s}_i}\backslash^{\square}_{\widetilde{s}_o}\backslash\square)^k.P_s$, if $s \in C_{i+1}$;

(g') $\overline{P}_s \triangleq \sum_{h\geq 0,k\geq 1}({}^{\overline{s}_i}\backslash^{\square}_{\overline{s}_o}\backslash\square)^h\, {}^{\underline{s}_i}\backslash^{\square}_{\underline{s}_o}\backslash\square({}^{\widetilde{s}_i}\backslash^{\square}_{\widetilde{s}_o}\backslash\square)^k.P_s$, if $s \notin C_{i+1}$

(h') $\overline{P}_s \triangleq \sum_{h\geq 0}({}^{\overline{s}_i}\backslash^{\square}_{\overline{s}_o}\backslash\square)^h {}^{\underline{s}_i}\backslash_{\underline{s}_o}.\overline{P}_s$, if $s \notin C_{i+1}$

where, by Lemma 12.2, h is the number of reactions requiring the absence of s plus possibly some reactions requiring s; and k is the number of reactions producing s.

It is worth nothing that, depending on the presence (P_s) or the absence ($\overline{P}_s$) of each entity s, for each process P_a (encoding a reaction a) the choice between the execution of the reaction code (points **(a)**, **(b)**, **(c)**) or the code expressing that reaction a is not applicable (point **(d)**) is deterministic. Also, the building of the code of process *Cxt* (points **(e)**, **(f)**), is univocally determined by the evolution of γ. It derives that the trend followed by the processes P_s (or $\overline{P}_s$) is also deterministic (points **(f)**, **(g)**, **(h)** or **(f')**, **(g')**, **(h')**), leading to $P' = [\![\mathcal{A}, \gamma^1]\!]$.

Corollary 17 (Correctness 2). *Let $P = [\![\mathcal{A}, \gamma]\!]$ and $j \geq 1$. If there exists P'' such that $P \xrightarrow{\tau\backslash_\tau \ldots {}^\tau\backslash_\tau}{}^{j} P''$, then letting $\pi^j = (\gamma^j, \delta^j)$ we have $P'' = [\![\mathcal{A}, \gamma^j]\!]$.*

Proof. We proceed by induction on the transition number $j \geq 0$.

base case $j = 1$: This case falls into the case of Proposition 16.

inductive case: We assume, by inductive hypothesis, that $\exists P'$ such that

$$P \xrightarrow{\tau\backslash_\tau \ldots {}^\tau\backslash_\tau}{}^{j-1} P'$$

and $P' = [\![\mathcal{A}, \gamma^{j-1}]\!]$. As P' is the encoding of an extended interactive process, by Proposition 16, it exists P'' such that $P' \xrightarrow{\tau\backslash_\tau \ldots {}^\tau\backslash_\tau} P''$, and $P'' = [\![\mathcal{A}, \gamma^j]\!]$.

Proposition 18 (Completeness 1). *Let $P = [\![\mathcal{A}, \gamma]\!]$ and $\pi^1 = (\gamma^1, \delta^1)$. Then, $P \xrightarrow{\tau\backslash_\tau \ldots {}^\tau\backslash_\tau} P' = [\![\mathcal{A}, \gamma^1]\!]$.*

PROOF. By Proposition 16, if there exists P' such that $P \xrightarrow{\tau\backslash_\tau \ldots \tau\backslash_\tau} P'$, then the structure of P' is deterministically computed.
Now, to prove that always exists P', we observe that even in the case no reaction a is applicable in the interactive process π in A, then process P can always execute a step transition, as its subprocesses P_a can always execute one of the *alternative code for when reaction a is not applicable* (see Definition 11, code for P_a processes).

Corollary 19 (Completeness 2). *Let* $P = [\![\mathcal{A}, \gamma]\!]$ *and* $\pi^j = (\gamma^j, \delta^j)$. *Then,* $P \xrightarrow{\tau\backslash_\tau \ldots \tau\backslash_\tau}{}^{j} P'' = [\![\mathcal{A}, \gamma^j]\!]$.

PROOF. The proof proceeds by induction on the number j, and it is similar to the one of Corollary 17.

Theorem 32 (Correspondence). $\sim_{\mathsf{F}} = \equiv_{\mathcal{L}_{\mathsf{F}}}$

PROOF. The proof is just an adaptation of the classical result. The two implications are proved separately.

$\sim_{\mathsf{F}} \subseteq \equiv_{\mathcal{L}_{\mathsf{F}}}$**:** Given any two processes $P \sim_{\mathsf{F}} Q$ we need to prove that for any bioHML formula G we have $P \models \mathsf{G}$ iff $Q \models \mathsf{G}$. Without loss of generality, we prove that $P \models \mathsf{G}$ implies $Q \models \mathsf{G}$. The proof is by structural induction on G.

- if $\mathsf{G} = \mathtt{t}$, then $Q \models \mathsf{G}$.
- if $\mathsf{G} = \mathtt{f}$, then the assumption $P \models \mathsf{G}$ is false and the implication holds.
- if $\mathsf{G} = \mathsf{G}_1 \wedge \mathsf{G}_2$ we take as inductive hypotheses that

$$\forall R, S.\ R \sim_{\mathsf{F}} S \wedge R \models \mathsf{G}_1 \Rightarrow S \models \mathsf{G}_1$$
$$\forall R, S.\ R \sim_{\mathsf{F}} S \wedge R \models \mathsf{G}_2 \Rightarrow S \models \mathsf{G}_2$$

  We need to prove that $Q \models \mathsf{G}$. Since $P \models \mathsf{G} = \mathsf{G}_1 \wedge \mathsf{G}_2$ we have $P \models \mathsf{G}_1$ and $P \models \mathsf{G}_2$. Since $P \sim_{\mathsf{F}} Q$, by inductive hypotheses we get $Q \models \mathsf{G}_1$ and $Q \models \mathsf{G}_2$. Hence $Q \models \mathsf{G}_1 \wedge \mathsf{G}_2 = \mathsf{G}$.
- if $\mathsf{G} = \mathsf{G}_1 \vee \mathsf{G}_2$ we take as inductive hypotheses that

$$\forall R, S.\ R \sim_{\mathsf{F}} S \wedge R \models \mathsf{G}_1 \Rightarrow S \models \mathsf{G}_1$$
$$\forall R, S.\ R \sim_{\mathsf{F}} S \wedge R \models \mathsf{G}_2 \Rightarrow S \models \mathsf{G}_2$$

  We need to prove that $Q \models \mathsf{G}$. Since $P \models \mathsf{G} = \mathsf{G}_1 \vee \mathsf{G}_2$ we have $P \models \mathsf{G}_1$ or $P \models \mathsf{G}_2$. If $P \models \mathsf{G}_1$, since $P \sim_{\mathsf{F}} Q$, by inductive hypotheses we get $Q \models \mathsf{G}_1$ and thus $Q \models \mathsf{G}_1 \vee \mathsf{G}_2 = \mathsf{G}$. If $P \models \mathsf{G}_2$, since $P \sim_{\mathsf{F}} Q$, by inductive hypotheses we get $Q \models \mathsf{G}_2$ and thus $Q \models \mathsf{G}_1 \vee \mathsf{G}_2 = \mathsf{G}$.

- if $\mathsf{G} = \langle\chi\rangle\mathsf{H}$ we take as inductive hypothesis that

$$\forall R, S.\ R \sim_{\mathsf{F}} S \wedge R \models \mathsf{H} \Rightarrow S \models \mathsf{H}$$

We need to prove that $Q \models \mathsf{G}$. Since $P \models \langle\chi\rangle\mathsf{H}$ it means that there exists υ, P' such that $P \stackrel{\upsilon}{\Rightarrow} P'$ with $\upsilon \models \chi$ and $P' \models \mathsf{H}$. Since $P \sim_{\mathsf{F}} Q$, there exists w, Q' such that $Q \stackrel{w}{\Rightarrow} Q'$ with $w \models \chi$ and $P' \sim_{\mathsf{F}} Q'$. Then, by inductive hypothesis, $Q' \models \mathsf{H}$ and thus $Q \models \langle\chi\rangle\mathsf{H} = \mathsf{G}$.

- if $\mathsf{G} = [\chi]\mathsf{H}$ we take as inductive hypothesis that

$$\forall R, S.\ R \sim_{\mathsf{F}} S \wedge R \models \mathsf{H} \Rightarrow S \models \mathsf{H}$$

We need to prove that $Q \models \mathsf{G}$. If there is no $\upsilon \models \chi$ such that $Q \stackrel{\upsilon}{\Rightarrow} Q'$ for some Q', then $Q \models [\chi]\mathsf{H} = \mathsf{G}$ trivially. For any υ, Q' such that $Q \stackrel{\upsilon}{\Rightarrow} Q'$ with $\upsilon \models \chi$, then as $P \sim_{\mathsf{F}} Q$ there must exist w, P' such that $P \stackrel{w}{\Rightarrow} P'$ with $w \models \chi$ and $P' \sim_{\mathsf{F}} Q'$. Since $P \models \mathsf{G} = [\chi]\mathsf{H}$ then it must be $P' \models \mathsf{H}$. Since $P' \sim_{\mathsf{F}} Q'$, by inductive hypothesis $Q' \models \mathsf{H}$. Hence $Q \models [\chi]\mathsf{H} = \mathsf{G}$.

$\equiv_{\mathcal{L}_{\mathsf{F}}} \subseteq \sim_{\mathsf{F}}$**:** We prove that $\equiv_{\mathcal{L}_{\mathsf{F}}}$ is a bio-simulation and thus included in $\sim_{\mathsf{F}}$. Take two generic processes $P \equiv_{\mathcal{L}_{\mathsf{F}}} Q$ and suppose $P \stackrel{\upsilon}{\Rightarrow} P'$ for some υ, P'.

- If $\upsilon \models \mathsf{F}$ we want to prove that there exists some w, Q' such that $Q \stackrel{w}{\Rightarrow} Q'$, with $w \models \mathsf{F}$ and $P' \equiv_{\mathcal{L}_{\mathsf{F}}} Q'$.

  Towards a contradiction, assume that we cannot find such w, Q'.

  If there is no transition $Q \stackrel{w}{\Rightarrow} Q'$ such that $w \models \mathsf{F}$, then the bioHML formula $\mathsf{G} \triangleq \langle\mathsf{F}\rangle\mathtt{t}$ is such that $P \models \mathsf{G}$ and $Q \not\models \mathsf{G}$, contradicting the assumption $P \equiv_{\mathcal{L}_{\mathsf{F}}} Q$.

  Otherwise, let $\mathcal{Q} \triangleq \{Q' \mid \exists w.\ Q \stackrel{w}{\Rightarrow} Q' \wedge w \models \mathsf{F}\}$ be the (non-empty) set of processes reachable from Q via a transition with a (complete) label that satisfies F. Since our processes are with guarded recursion, the set $\mathcal{Q}$ is finite. Let $\mathcal{Q} = \{Q'_1, ..., Q'_n\}$. By hypothesis all processes in $\mathcal{Q}$ must not be bio-logically equivalent to P', hence for any $i \in [1, n]$ there exists a bioHML formula G_i such that $P' \models \mathsf{G}_i$ and $Q'_i \not\models \mathsf{G}_i$ (if it was the opposite, $P' \not\models \mathsf{H}_i$ and $Q'_i \models \mathsf{H}_i$ for some H_i, we can use the converse formula $\mathsf{G}_i \triangleq \overline{\mathsf{H}}_i$). But then the formula $\mathsf{G} \triangleq \langle\mathsf{F}\rangle(\mathsf{G}_1 \wedge \cdots \wedge \mathsf{G}_n)$ is such that $P \models \mathsf{G}$ and $Q \not\models \mathsf{G}$, contradicting the assumption $P \equiv_{\mathcal{L}_{\mathsf{F}}} Q$.

- If $\upsilon \not\models \mathsf{F}$ then the proof is analogous to the previous case (by exploiting $\neg\mathsf{F}$) and thus omitted.