

DYNAMIC SYNC-PROGRAMS FOR MODULAR VERIFICATION OF BIOLOGICAL SYSTEMS

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Abstract

We propose dynamic sync-programs, a bio-inspired automata-based formalism for the description and the modular verification of properties of biological systems. The formalism allows entities to be created dynamically, in particular by other already running entities, as it often happens in biological systems. Moreover, multiple copies of the same entities can be present at the same time in a system. Modular verification allows properties expressed as DACTL formulae (dynamic version of the universal fragment of CTL) to be verified on a portion of the model, rather than on the whole model. We show, as an example, the application of our approach to a biological case study, namely the EGF signalling pathway.

1. Introduction

Model checking permits the verification of properties of a system (expressed as logical formulae) by exploring all the possible behaviours of the system. Recently, it has been successfully applied to analysis of biological systems (see e.g. [7, 5, 2]). However, model checking techniques have traditionally suffered from the state explosion problem. A method for trying to avoid such a problem is to use the natural decomposition of the system. The goal is to verify properties of individual components and infer that these hold in the complete system. A technique proposed by Attie [1] exploits the preservation of properties expressed with ACTL (the universal fragment of the CTL temporal logic) from components to whole systems in order to verify concurrent programs and synthesise systems from specifications. Attie uses a formalism called *synchronisation skeletons* [3] suitable for describing distributed systems.

In [4] we proposed an extension of Attie's approach for the description and the modular verification of biological systems. In particular, we defined *sync-programs*, an automata-based formalism of interactive systems which extends the formalism used by Attie by allowing processes to perform transitions simultaneously. Here we further extend the approach by allowing sync-automata (the components of a sync-program) to be created dynamically by other already running sync-automata. Moreover, we allow several instances of the same sync-automata to be

executed concurrently, without any bound of the number of concurrent instances of the same sync-automaton. As our previous extension of Attie’s approach, also this further extension is biologically motivated. Indeed, in biological systems it is often the case that entities involved in the processes of interest (e.g. proteins) are synthesised by other active entities (e.g. the DNA). Moreover, it is very common that a number of copies of the same entities (e.g. proteins) are active at the same time.

In what follows we give a formal definition of the syntax of dynamic sync-programs and their translation into Petri nets. The existence of such a translation implies that the formalism is not Turing-complete and that verification of the properties of interest is decidable. In order to shorten the presentation, we define the semantics of dynamic sync-programs by exploiting the translation into Petri nets and their semantics. Subsequently, we develop the modular verification approach for dynamic sync-programs by defining projection operations on their syntax and on their semantics, and by giving some results on the preservation of properties from components to whole systems. Finally, we apply our approach to a biological case study, namely the EGF signalling pathway.

2. Dynamic sync-programs

2.1. Syntax

To model biological systems, we use a component-based approach. Each component represents a biological entity, e.g. a protein or an enzyme. The system will be modelled by a program consisting of a parallel composition of automata of different types. We assume a finite *set of component types* CT .

For the purpose of developing a modular approach, the interactions between component types are indicated explicitly. We say that (not necessarily distinct) types i and j from CT *interact directly*, when one can perform an activity conditioned on the activity of the other and vice versa or when component of one type causes creation of a component of the other type. Direct interaction is distinguished from indirect interaction, which is mediated by a third party. We define an undirected *interaction graph* I (see e.g. Figure 1), where the nodes are types from CT and there is an edge between nodes i and j iff types i and j interact directly. Note that there can be loops in the graph, representing interaction of two components of the same type. We use notation iIj if there is an edge between i and j in graph I . By $|I|$ we denote the set of nodes in I and by $I(i)$ the set $\{j \in |I| \mid iIj\}$. By $J \subseteq I$ we denote a connected subgraph J of I .

In our approach, states of the components will be encoded by using atomic propositions. With every component type i a set AP_i of *atomic propositions* is associated. The sets of atomic propositions are pairwise disjoint for all the types, i.e. if $i \neq j$ then $AP_i \cap AP_j = \emptyset$. We assume a function $type : AP \rightarrow CT$ that for an atomic proposition from AP_i gives its type i . Analogously, function $types$ yields a multiset of types of a set of APs.

An individual component of a component type from CT is modelled by a finite state machine called sync-automaton.

Definition 1. A *sync-automaton* A_i^I , where i is a component type and I is an interaction graph, is a tuple $(S_i, S_i^0, SC_i, CC_i, R_i)$:

- $S_i \subseteq \mathcal{P}(AP_i)$ is the set of *states*
- $S_i^0 \subseteq S_i$ is the set of *initial states*
- SC_i is a set of labels of the form $\bigwedge_{k \in L} U_k : V_k$ where finite $L \subseteq \mathbb{N}$ and for every $k \in L$ there is a $j \in I(i)$ such that U_k, V_k are sets of atomic propositions drawn from AP_j or their negations. A label from SC_i is called a *synchronisation condition*
- CC_i is a set of labels of the form $new(\bigwedge_{k' \in K} W_{k'})$ where finite $K \subseteq \mathbb{N}$ and for every $k' \in K$ there is a $j \in I(i)$ such that $W_{k'}$ is a set of atomic propositions drawn from AP_j or their negations. A label from CC_i is called a *creation condition*
- $R_i \subseteq S_i \times SC_i \times CC_i \times S_i$ are the *moves* between states.

Each state of a sync-automaton A_i^I is a truth value assignment to atomic propositions of component type i . Each move is labelled by a synchronisation condition sc and by a creation condition cc . We denote a move from state s_i to state t_i with labels sc and cc by $s_i \xrightarrow{sc, cc} t_i$. This move intuitively means that automaton A_i^I can move from s_i to t_i if there are concurrently performed moves of sync-automata of types in $I(i)$ satisfying condition sc . By performing the move, automata described by cc are created.

An example of a sync-automaton can be seen on Figure 2. The component type uses two atomic propositions, A and B . The two states of the automaton are $\{A, B\}$ and $\{\neg A, B\}$ with the former chosen to be initial. Moves between states are labelled by synchronisation and creation conditions.

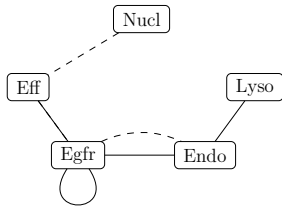


Figure 1: The interaction graph of the model of the EGF pathway (dashed lines – creation).

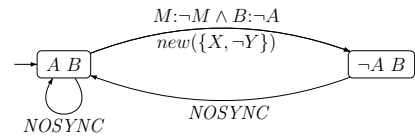


Figure 2: Example of a sync-automaton and move conditions.

The synchronisation condition is a label in form $\bigwedge_{k \in L} U_k : V_k$ and specifies requirements for automata of types from $I(i)$ to synchronise with. For every k in L , the sets of propositions U_k and V_k are to be satisfied in the starting and ending state, respectively, of the concurrently performed move of a sync-automaton of a type $type(U_k)$ (see Figure 2).

We remark that in the synchronisation condition of a move of a sync-automaton of type i there can be multiple references to the components of a type j , referring to different instances

of sync-automata of such a type. References to other instances of the same type i are also allowed. Moreover, note that it is possible for L to be empty. Intuitively, this means that the sync-automaton A_i^I does not have any requirements on other sync-automata. We write a synchronisation condition of this form, i.e. $\bigwedge_{j \in \emptyset} U_k \cdot V_k$, as *NOSYNC*. Move $s_i \xrightarrow{\text{NOSYNC}} t_i$ represents an autonomous move of A_i^I .

The creation condition is a label in form $\text{new}(\bigwedge_{k' \in K} W_{k'})$ that specifies sync-automata that are to be created. For every k' in K , the set of atomic propositions (or their negations) $W_{k'}$ encodes an initial state of sync-automaton A_j^I for $j = \text{type}(W_{k'})$ which will be created having this initial state. In case the multiset K is empty, the creation condition is not displayed. For an example see the move on Figure 2 from $\{A, B\}$ to $\{\neg A, B\}$ which specifies a creation of an automaton with initial state $\{X, \neg Y\}$.

By running in parallel sync-automata of types related by an interaction graph, we obtain a sync-program. Note that a sync-program can contain none or multiple sync-automata of any component type.

Definition 2. Let I be an interaction graph. A *sync-program* is a tuple $P^I = (s_{i_1} || \dots || s_{i_n})$, where each s_i is an initial state of sync-automaton A_j^I of type $j = \text{type}(s_i)$ where $j \in I$. The creation conditions on the moves of all sync-automata are well formed, i.e. for every creation condition $\text{new}(\bigwedge_{k' \in K} W_{k'})$, each $W_{k'}$ describes a unique state s' of a sync-automaton of type in I and s' is initial.

2.2. Sync-programs as Petri nets

A dynamic sync-program can be represented as a Petri net. Intuitively, a place in the net represents a state of a component type, a transition embodies synchronisation of sync-automata and creation of new ones. Then number of tokens in a place represents the number of automata currently in that state. A Petri net consists of a Petri net graph and an initial marking. Note that the net graph is created from the set of all component types and is always finite. Then the initial marking is constructed according to the program.

Definition 3. Given a set of sync-automata $A_I = \{A_i \mid i \in |I|\}$, we construct a *Petri net graph* $G_I = (S_I, T_I, W_I)$ as follows. The set of places is $S_I = \bigcup_{i \in |I|} S_i$. We assume a linear order on S_I . The set of transitions T_I is such that $T_I \subseteq S_I^* \times S_I^*$ where $S_I^* = \bigcup_{k=1}^{\infty} S_I^k$, with $S_I^1 = S_I$ and $S_I^k = S_I \times S_I^{k-1}$. Set T_I contains transitions of the form $tr = (In, Out)$ with In ordered. We denote with In_i and Out_i the i -th element of In and Out , respectively. Moreover, $(In, Out) \in T_I$ if and only if In and Out are minimal tuples and $Create \subseteq \mathbb{N} \times \mathcal{P}(AP)$ is a minimal set satisfying:

1. for all $i \in \{1, \dots, |In|\}$ there is a move $In_i \xrightarrow[\text{new}(\bigwedge_{k' \in K} W_{k'})]{\bigwedge_{k \in L} U_k \cdot V_k} Out_i$ in $A_{\text{type}(In_i)}$ such that
 - there is an injective mapping $inst$ from L to $\{1, \dots, |In|\} - \{i\}$ such that for all $k \in L$ for all $p \in U_k$: $In_{inst(k)}(p) = tt$ and for all $p \in V_k$: $Out_{inst(k)} = tt$

- $\{(k', W_{k'}) \mid k' \in K\} \subseteq Create$
- 2. for all $i \in \{1, \dots, |Create|\}$: $Out_{|In|+i} = c_i$ where $Create_i = (j_i, c_i)$

The set of arcs $W_I \subseteq (S_I \times T_I) \cup (T_I \times S_I)$, consists of the union of the following sets

- $\{(s, |Move|, tr) \mid tr \in T_I \wedge tr = (In, Out) \wedge Move = \{j \mid (s, j) \in In\} \wedge Move \neq \emptyset\}$
- $\{(tr, |Move|, t) \mid tr \in T_I \wedge tr = (In, Out) \wedge Move = \{j \mid (t, j) \in Out\} \wedge Move \neq \emptyset\}$.

In the definition of the encoding of dynamic sync-programs into Petri net graphs we have represented a transition as a pair of tuples (In, Out) . Tuple In consists of the starting states of the moves that synchronise in the step represented by the transition. The first $|In|$ elements of tuple Out represent the ending states reached by such moves from the corresponding elements in In . The rest of tuple Out represents the initial states of the sync-automata dynamically created in this step. The set $Create$ is used to collect such initial states from the creation conditions of the moves involved in the transition. Minimality of In, Out and $Create$ ensures the indivisibility of the synchronisation, and guarantees that all the considered synchronising moves can be grouped in a single transition of the Petri net.

With this encoding we have that Petri nets can be considered as an alternative representation for sync-programs. Petri nets are very intuitive and allow the modeller to have a comprehensive view of the system. However, since they are not compositional, they are not suitable neither for modular modelling, nor for modular verification. The existence of the translation of sync-programs into Petri nets is very important since it implies that the formalism is not Turing-complete and that verification of the properties of interest is decidable.

2.3. Semantics

Now we exploit the translation of dynamic sync-programs into Petri nets to give a formal semantics to such programs. The semantics could also be defined directly on the sync-programs (as in the non-dynamic case in [4]). However, we choose not to follow the latter approach here for reasons of space limitation. The semantics of a Petri nets is given in terms of a transition system. Since we consider only nets obtained from the encoding of sync-programs, we can give a slightly richer transition system that includes some information about the sync-automata involved in each transition.

A *marking* of a Petri net graph is a multiset of its places, i.e. a mapping $M : S_I \rightarrow \mathbb{N}$. We say that a marking assigns a number of tokens to each place. Note that markings can be added like any multiset: $M + M' = \{s \rightarrow (M(s) + M'(s)) \mid s \in S_I\}$. The *execution* of a Petri net graph $G_I = (S_I, T_I, W_I)$ can be defined as the transition relation \rightarrow^l on its markings, as follows:

- for any t in T_I : $M \rightarrow_t^l M'$ iff there is $M'' : S_I \rightarrow \mathbb{N}$ s.t. $M = M'' + \sum_{s \in S_I} W_I(s, t) \wedge M' = M'' + \sum_{s \in S_I} W_I(t, s)$ and $l = types(t)$

- $M \rightarrow_l M'$ iff there is $t \in T_I$ s.t. $M \xrightarrow{t} M'$

In words: firing a transition t in a marking M consumes $W_I(s, t)$ tokens from each of its input of its input places s , and produces $W_I(t, s)$ tokens in each of its output places s and label l contains types of automata that participate in the transition. These types are recorded for the purpose of compatibility of the semantics obtained through the translation in Petri nets and the direct semantics where they are necessary.

We say that a marking M' is reachable from a marking M in one step if $M \rightarrow_l M'$ for some l . We say that it is *reachable* from M if $M \xrightarrow{*} M'$, where $\xrightarrow{*}$ is the transitive closure of \rightarrow_l , that is, if it is reachable in zero or more steps.

A *Petri net* consists of a Petri net graph and of an initial marking M_0 . The initial marking of a Petri net is obtained from the encoding of a sync-program. It contains as many tokens in a place representing an initial state of a sync-automaton from I , as there are instances of such initial state in program P^I . For a Petri net $N_I = (S_I, T_I, W_I, M_0)$ the set of reachable markings is the set $R(N_I) = \{M' \mid M_0 \xrightarrow{*} M'\}$. The *reachability graph* of N_I is the transition relation \rightarrow_l restricted to $R(N)$.

Definition 4 (Semantics). The *I-structure* \mathcal{M}_I of a dynamic I -program P^I is defined as the reachability graph of N_I , where N_I is the marked Petri net obtained by the encoding of P^I . A transition (M, l, M') is in \mathcal{M}_I iff $M \rightarrow_l M'$ is in $R(N_I)$. A marking from \mathcal{M}_I is called an *I-state*. The set of initial sets of \mathcal{M}_I is a set of initial markings that is admissible by all automata types.

A concept that will allow us to reason about properties of programs is that of path. A *path* in an I -structure \mathcal{M}_I is a sequence of I -states and transition labels $\pi = (s^1, l^1, s^2, l^2, \dots)$ such that for all m , $(s^m, l^m, s^{m+1}) \in \mathcal{M}_I$. A *fullpath* is a maximal path, and it is infinite unless for some $s^{m'}$ there is no $s^{m'+1}$ and $l^{m'}$ such that $(s^{m'}, l^{m'}, s^{m'+1}) \in \mathcal{M}_I$. Let π^m denote the suffix of π starting in m -th I -state.

3. Subprograms and projections

In order to develop our modular verification techniques we need two projection operations that allow us to restrict the syntax and the semantics of a sync-program to the components that have an influence on the satisfaction of a property of interest.

A *sync-subprogram* represents the behaviour of a portion of a sync-program in isolation. We obtain a sync-subprogram by projecting a sync-program P^I onto an interaction graph $J \subseteq I$. Graph J represents a subset (subgraph) of the set of all component types I . We denote the projection by the projection operator $\downarrow J$. In order to obtain $P^I \downarrow J$ we compute a $J' \supseteq J$ which is a subgraph of I containing types of automata that can create (also by transitivity) automata of types in J . Then by projection $P^I \downarrow J'$ inspired by [4] we obtain the *syntactical projection* $P^I \downarrow J$.

Now, let J' be the graph that can be obtained through syntactical analysis of the component types as follows. The *creation graph of J* is a graph with vertices from CT . There is an oriented edge from i to j if there is a move in A_i^I whose creation condition contains an initial state of automaton of type j . Subgraph J' is induced by vertices from J together with vertices from whose there is a path to vertices in J . Now denote $X := J$, $Y := J' - J$, $Z := I - J'$. The projection onto J' removes all automata from Z , namely those that neither are nor create (not even by transitivity) an automaton from J . This is defined as follows.

Definition 5. Let $J' \subseteq I$ be an interaction graph, where $|J'| = \{j_1, \dots, j_k\}$. Let $P^I = (s_{i_1} || \dots || s_{i_n})$ with $A_i^I = (S_i, S_i^0, SC_i, CC_i, R_i)$ and $s_i \in S_i^0$ for each $i \in |I|$. Then $P^I \downarrow J' = (s_{j_1} || \dots || s_{j_k})$ with $A_j^J = (S_j, S_j^0, SC'_j, CC'_j, R'_j)$ and $s_j \in S_j^0$ for each $j \in |J|$ where:

- $SC'_j = \{\wedge_{j' \in L \cap |J'|} U_{j'}:V_{j'} \mid \wedge_{j' \in L} U_{j'}:V_{j'} \in SC_j\}$;
- $CC'_j = \{new(\wedge_{j' \in K \cap |J'|} W_{j'}) \mid new(\wedge_{j' \in K} W_{j'}) \in CC_j\}$;
- $R'_j = \{s_j \xrightarrow[\text{new}(\wedge_{j' \in K \cap |J'|} W_{j'})]{\wedge_{j' \in L \cap |J'|} U_{j'}:V_{j'}} t_j \mid s_i \xrightarrow[\text{new}(\wedge_{j' \in K} W_{j'})]{\wedge_{j' \in L} U_{j'}:V_{j'}} t_j \in R_j\}$.

The projection contains only sync-automata from J' . Each sync-automaton has the same states as its counterpart in P^I but the synchronisation and creation conditions on their moves concern only sync-automata from J' . We remark that a sync-subprogram $P^I \downarrow J'$ is still a sync-program with interaction graph J' , hence it can be also denoted by $P^{J'}$. Thus, syntactical projection is defined as follows: $P^I \downarrow J = P^I \downarrow J'$ where J' is obtained from the creation graph of J .

To define the semantic projection, let us denote with $s \upharpoonright J$ the projection of an I -state s onto a $J \subseteq I$. It is the greatest submultiset of s consisting only of states of sync-automata from J . A transition (s, l, t) in a I -structure can be projected onto $J \subseteq I$ as follows: $(s, l, t) \upharpoonright J = (s \upharpoonright J, l \cap |J|, t \upharpoonright J)$. Projection of a I -structure \mathcal{M}_I onto J can now be defined as the transition relation consisting of the projections onto J of the transitions in \mathcal{M}_I . Note that it is possible that the transition label is an empty set, and the corresponding projected states can be both equal (synchronisation outside J was removed) or different (because of creation by an automaton outside J). However this does not cause problems, as for the purpose of verification the transitions of the former type are removed.

Path projection $\pi \upharpoonright J$ is obtained by projecting every transition of the path π onto J .

4. Modular verification

In order to analyse the behaviour of a biological system we would like to verify properties of computations of sync-program P^I representing the system. Assume, that property ϕ_J only regards part of the system, in particular the part involving only components from $J \subseteq I$. We would like to check satisfaction of ϕ_J on semantics of P^I . In order to avoid the space explosion, we want to check it on a smaller and more abstract semantics, in particular semantics of $P^J = P^I \downarrow J$. By abstracting away some of the automata we lose some synchronisations,

obtaining an overapproximation of the behaviour of P^I . The essence of the verification can be expressed as equation $Sem \circ \lceil \subseteq \lceil \circ Sem$, where Sem denotes the semantic function.

In this section we assume that sync-programs contain no indirect synchronisations in the form of synchronisation conditions. If this is not the case, the program must be modified in a preprocessing step. We argue, that each computation being a fair path of the entire system projected onto J is present as a computation of P^J . In the line of [4] we first prove the Transition projection and Path projection lemmas. By assuming fairness we avoid problems of losing the eventuality properties and we get the Fullpath projection lemma.

Subsequently, we specify the logic suitable for describing the properties of dynamical sync-programs and finally we show the preservation of this logic.

4.1. Lemmas

Lemma 4.1 (Transition projection). *Let I be an interaction graph and \mathcal{M}_I the semantics of sync-program P^I . For all I -states s, t in \mathcal{M}_I and all multisets l of elements from $|I|$, transition (s, l, t) is in \mathcal{M}_I iff for all $J \subseteq I$ we have that $(s, l, t) \lceil J$ is in \mathcal{M}_J , where \mathcal{M}_J is the semantics of sync-program $P^J = P^I \lceil J$.*

This lemma can be proved by using the definition of semantics and the definitions of both projections.

The path projection result follows from using the previous lemma for every transition in the path.

Lemma 4.2 (Path projection). *Let I be an interaction graph and \mathcal{M}_I semantics of sync-program P^I . For every $J \subseteq I$ if π is a path in \mathcal{M}_I then $\pi \lceil J$ is a path in \mathcal{M}_J , where \mathcal{M}_J is the semantics of sync-program $P^J = P^I \lceil J$.*

By assuming fairness of paths we immediately get the required Fullpath projection lemma.

Definition 6. A path $\pi = (s^1, l^1, s^2, l^2, \dots)$ in \mathcal{M}_I is *fair* iff for all $i \in |I|$ we have that $\{m \mid i \in l^m\}$ is infinite.

Lemma 4.3 (Fullpath projection). *Let $J \subseteq I$ be an interaction graph. If π is a fair fullpath in \mathcal{M}_I , then $\pi \lceil J$ is a fair fullpath in \mathcal{M}_J .*

4.2. Logic

We define a logic for specification of properties of dynamic sync-programs. We adapt the logic ACTL [6], the universal fragment of CTL, for our purpose. We need to incorporate two aspects, namely dynamicity since automata can be added in runtime, and reasoning about instances since there can be more automata of a given type.

Definition 7. *Syntax of DACTL* is defined inductively as follows:

- The constants *true* and *false* are formulae.
- Program formula is a formula
 - Program formula is of the form $pf = pc_1 \vee \dots \vee pc_n$ where pc_j is a program conjunction for all j
 - Program conjunction is of the form $pc_j = tf[i_1] \wedge \dots \wedge tf[i_{|CT|}]$ and $tf[i]$ is a type formula for all i
 - Type formula is of one of the forms
 - * $tf[i] = if(k_1) \wedge \dots \wedge if(k_t)$ where type index $i \in CT$ and $if(k)$ is an instance formula for all k
 - * $\otimes i$ where $i \in CT$
 - * $\otimes if(1)$
 - Instance formula is of the form $if(k) = p_1 \wedge \dots \wedge p_{n_1}$ where instance index $k \in \mathbb{N}$ and the atomic proposition $p_m \in AP_{type(k)}$ or its negation for all m
- If f, g, h are formulae, then so are $f \wedge A[gUh]$ and $f \vee A[gUh]$.
- If f, g, h are formulae, then so are $f \wedge A[gU_w h]$ and $f \vee A[gU_w h]$.

We define the logic $DACTL_J$ to be DACTL where the instance formulae are drawn from $AP_J = \bigcup_{j \in |J|} AP_j$. Abbreviations in DACTL: $AFf \equiv A[true U f]$ and $AGf \equiv A[f U_w false]$.

In this logic, state formulae called *program formulae* can describe states of multiple automata of the same type. We are not able to reference a particular automaton amongst the automata of a given type, but we still can specify different copies of the type. To be able to distinguish which automaton the property refers to, atomic propositions are labelled by distinct indices. For example property $a(1) \wedge a(2)$ talks about two automata of the same type, and both of them need to be in a state that satisfies a . Thus also $a(1) \wedge \neg a(2)$ is a consistent property. For convenience and intuitiveness the program formulae are in the DNF.

Properties expressible by DACTL formulae are similar to ACTL properties and thus are able to represent a significant class of biologically relevant properties [4].

Definition of the semantics of DACTL formulae on the I -structure follows. Note that only fair fullpaths are considered.

Definition 8. *Semantics of DACTL.* We define $\mathcal{M}_I, s \models f$ (resp. $\mathcal{M}_I, \pi \models f$) meaning that f is true in structure \mathcal{M}_I at state s (resp fair fullpath π). We define \models inductively:

- $\mathcal{M}_I, s^1 \models true$. $\mathcal{M}_I, s^1 \not\models false$.
- $\mathcal{M}_I, s^1 \models pf$ iff $pf = pc_1 \vee \dots \vee pc_n$ and there is $n' \in \{1, \dots, n\}$ such that $\mathcal{M}_I, s^1 \models pc_{n'}$

- $\mathcal{M}_I, s^1 \models pc_j$ iff $pc_j = tf[i_1] \wedge \dots \wedge tf[i_{|CT|}]$ and for all $i \in \{i_1, \dots, i_n\} \subseteq CT$:
 - $\mathcal{M}_I, s^1 \models tf[i]$
 - $\mathcal{M}_I, s^1 \models tf_i$ iff
 - * $tf[i] = if(k_1) \wedge \dots \wedge if(k_t)$ and there is a mapping $inst$ from $\{k_1, \dots, k_t\}$ to states of S_i such that the multiset $\{inst(k_1), \dots, inst(k_t)\} \subseteq s^1 \upharpoonright i$ and for all $k \in \{k_1, \dots, k_t\}$ holds $\mathcal{M}_I, inst(k) \models if(k)$
 - * $tf[i] = \otimes i$ and $s^1 \upharpoonright i = \emptyset$
 - * $tf[i] = \otimes if(1)$ and for all s' in $s^1 \upharpoonright i$: $\mathcal{M}_I, s' \models if(1)$
 - $\mathcal{M}_I, s' \models if[k]$ iff $if(k) = p_1 \wedge \dots \wedge p_{n_k}$ and for all $m \in \{1, \dots, n_k\}$ holds $\mathcal{M}_I, s' \models p_m$
- $\mathcal{M}_I, s^1 \models f \wedge A[gUh]$ iff $\mathcal{M}_I, s^1 \models f$ and for every fair fullpath $\pi = (s^1, l^1, \dots)$ in \mathcal{M}_I : there exists $m \in \mathbb{N}$ such that $\mathcal{M}_I, \pi^m \models h$ and for all $m' < m$: $\mathcal{M}_I, \pi^{m'} \models g$. The \vee case is analogous.
- $\mathcal{M}_I, s^1 \models f \wedge A[gU_w h]$ iff $\mathcal{M}_I, s^1 \models f$ and for every fair fullpath $\pi = (s^1, l^1, \dots)$ in \mathcal{M}_I : for all $m \in \mathbb{N}$, if for all $m' < m$: $\mathcal{M}_I, \pi^{m'} \not\models h$, then $\mathcal{M}_I, \pi^m \models g$. The \vee case is analogous.

As an example of satisfaction, consider $AP_{t_1} = \{a, b\}$ and $AP_{t_2} = \{X, Y\}$. Denote the state $\{a \wedge b, a \wedge \neg b\}$ as s . Now we have that $s \models b(1) \wedge \neg b(2)$, $s \models \neg b(1) \wedge b(2)$, but $s \not\models \neg b(1) \wedge \neg b(2)$ and $s \not\models b(1) \wedge \neg b(2) \wedge b(3)$. Special state operator \otimes talks about nonexistence of an automaton of a given type. For example $s \models \otimes t_2$ but $s \not\models \otimes t_1$. On the other hand \otimes says that an instance formula holds in all the instances of a given type, for example $s \models \otimes a$ but $s \not\models \otimes b$.

4.3. Property preservation theorem

The following theorem guarantees that successful verification of a $DACTL_J$ property in M_J amounts to its successful verification in M_I .

Theorem 4.4 (Property preservation). *Let $J \subseteq I$ be an interaction graph, s an I -state and f a $DACTL_J$ property. If $\mathcal{M}_J, s \upharpoonright J \models f$ then $\mathcal{M}_I, s \models f$.*

The theorem can be proved by induction on the structure of the formula. The base case holds because of the fact that the sets of atomic propositions are pairwise disjoint and the J -state is maintained by the semantic projection. The inductive cases are proved with the help of the Fullpath projection lemma.

5. An Application: the EGF Signalling Pathway

In Biology, signal transduction refers to any process by which a cell converts one kind of signal into another. Signals are typically proteins that may be present in the environment of the cell. In order to recognise that a signal is available in the environment, a cell exposes some receptors on its external membrane. A complex signal transduction cascade that modulates cell proliferation is based on a family of receptors called epidermal growth factor receptors (EGFRs)

that are produced by specific genes in the DNA (through the RNA) and are located on the cell surface. Receptors are activated by the binding with a specific ligand (epidermal growth factor, EGF) to form a ligand-receptor complex. After activation, EGFR undergoes a transition from a monomeric form to a dimeric one (consisting of two ligand-receptor complexes). EGFR dimerisation enables activation of effector proteins, which initiates several signal transduction cascades, leading to cell proliferation. Occasionally, ligand-receptor dimers are internalised in endosomes and consequently targeted for degradation inside the lysosomes.

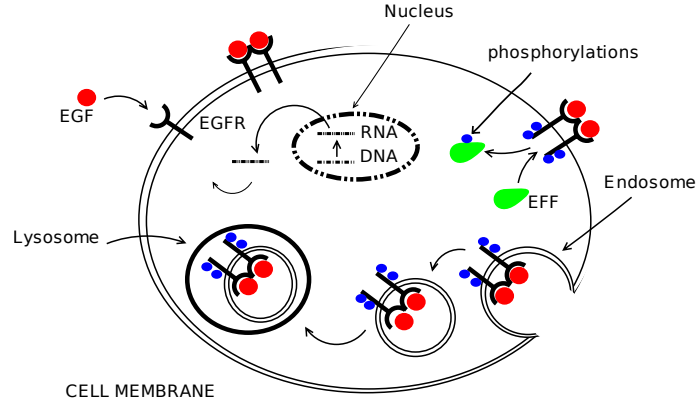


Figure 3: The first steps of the EGF pathway.

We model the described steps of the EGF pathway (with some simplifications) as a dynamic sync-program. The set of component types consists of five types: type $nucl$, with $AP_{nucl} = \{Dna\}$; type $egfr$ with $AP_{egfr} = \{Egfr, On_membr, Bound, Dim, Cr_endo\}$; type eff with $AP_{eff} = \{Eff_p\}$; type $endo$, with $AP_{endo} = \{Endo, In_lyso\}$; and type $lyso$, with $AP_{lyso} = \{Lyso\}$. The interaction graph I of the model is shown in Figure 1.

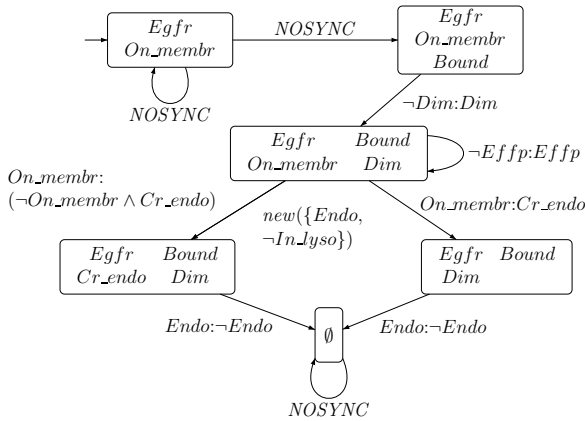


Figure 4: A_{egfr} .

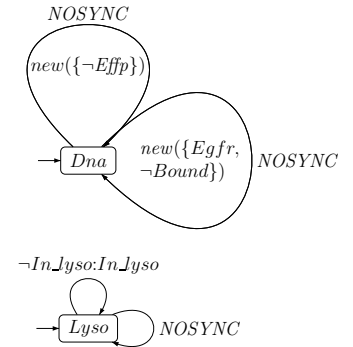
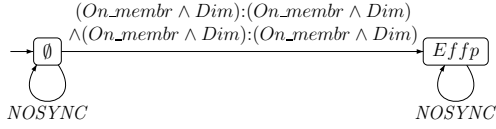
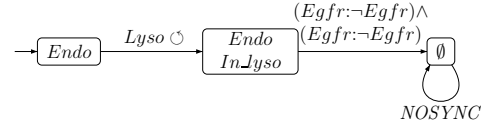


Figure 5: A_{nucl} and A_{lyso} .

Sync-automaton A_{nucl} (depicted in Figure 5) describes the DNA/RNA activity in the nucleus, that is the production of receptors and effectors. Sync-automaton A_{egfr} (Figure 4) describes a receptor. It starts by binding an EGF signal, as modelled by the first $NOSYNC$ move. (Absence of EGF signals in the environment is modelled by the self-looping $NOSYNC$ move

in the initial state.) Subsequently, it synchronises with another sync-automaton of the same type to form a dimer (represented by two automata in a state in which Dim holds). The dimer can then interact with effectors. A dimer can be internalised by an endosome. The endosome is created by an automaton in a state in which Dim holds, by synchronising with another automaton in the same state (assumed to be the other component of the dimer). In order to ensure that only one endosome is created, the moves of the two components of the dimer are kept distinct. Finally, a dimer synchronises with an endosome in order to be destroyed (inside the lysosome).

Figure 6: A_{eff} .Figure 7: A_{endo} .

An effector protein is modelled by sync-automaton A_{eff} (Figure 6) that, once created, can either do nothing (the $NOSYNC$ self-looping move) or synchronise with a dimer in order to move to a state in which it is phosphorylated. Finally, sync-automaton A_{endo} (Figure 7) describes an endosome that synchronises first with a sync-automaton A_{lyso} (Figure 5) to model entrance into the lysosome and then with two automata assumed to represent the dimer it is carrying.

The sync-program corresponding to the initial configuration of the model is $P^I = (Dna || Lyso)$. Now, we discuss properties that could be verified on the model in a modular way. We first show some properties about the correctness of the model itself:

- “The number of instances of A_{egfr} in which Dim holds is always even“. Since we cannot reason on the numbers of instances, what we can actually prove is the following weaker property: “In every reachable state Dim holds in either zero or at least two instances of A_{egfr} ”. This property is expressed by the formula $AG(\otimes Egfr \vee \otimes \neg Dim \vee (Dim(1) \wedge Dim(2)))$ that can be verified on the semantics of P^{egfr} .
- “If an endosome is destroyed, then at the same time two receptors are destroyed”. This property is expressed by the formula $AG(\otimes Endo U_w AG(Endo(1) \wedge Egfr(2) \wedge Egfr(3) U \neg Endo(1) \wedge \neg Egfr(2) \wedge \neg Egfr(3)))$ that can be verified on the semantics of $P^{endo, egfr}$.

Moreover, some examples of properties of the behaviour of the modelled system are the following:

- “If EGF signals are present in the environment, eventually either effectors are phosphorylated or endosomes are created.” is a property stating the correct functioning of the model. It is expressed as the formulae: $AG(\otimes Egfr \vee \neg Bound U_w AF(Effp \vee Endo))$ that can be proved to hold on the semantics of $P^{egfr, eff, endo}$.

- “If an endosome is created, it will be eventually destroyed”. Actually what we can prove is the weaker property stating “If an endosome is created, an endosome will be eventually destroyed”. This property is expressed by the formula $\otimes Endo U_w AF \neg Endo$ that can be verified on the semantics of P^{endo} .

6. Discussion

We have proposed a dynamic automata-based formalism for the description and modular verification of biological systems. The proposed approach extends our previous proposal [4] with the possibility of dynamically creating new instances of automata. This made the formalism much more suitable to describe biological systems, but required the modular verification approach to be adapted. Indeed, in this case verification of a property cannot be performed only on the components of the system that are directly involved in the property, but also on those that may cause (even indirectly) the creation of some components involved in the property. Hence, we introduced a new syntactic projection operation that takes these aspects into account. Such an operation could be improved (i) by removing unnecessary *NOSYNC* moves in the components not directly involved in the property, and (ii) by removing synchronisations between components directly and non-directly involved in the property. Verification of DACTL properties can be limited to a portion of the model, which renders the model checking more efficient. However, precise evaluation of obtained reduction in terms of state space and time is left for further investigation.

References

- [1] Paul C. Attie. Synthesis of large dynamic concurrent programs from dynamic specifications. *CoRR*, abs/0801.1687, 2008.
- [2] Federica Ciocchetta and Jane Hillston. Bio-pepa: A framework for the modelling and analysis of biological systems. *Theor. Comput. Sci.*, 410(33-34):3065–3084, 2009.
- [3] Edmund M. Clarke and E. Allen Emerson. Design and synthesis of synchronization skeletons using branching-time temporal logic. In *Logic of Programs, Workshop*, pages 52–71, London, UK, 1982. Springer-Verlag.
- [4] Peter Drábik, Andrea Maggiolo-Schettini, and Paolo Milazzo. Modular verification of interactive systems with application to biology. *CS2Bio’10. In press*, 2010.
- [5] François Fages, Sylvain Soliman, and Nathalie Chabrier-Rivier. Modelling and querying interaction networks in the biochemical abstract machine biocham. *Journal of Biological Physics and Chemistry*, 4:64–73, 2004.
- [6] Orna Grumberg and David E. Long. Model checking and modular verification. *ACM Trans. Program. Lang. Syst.*, 16(3):843–871, 1994.
- [7] John Heath, Marta Kwiatkowska, Gethin Norman, David Parker, and Oksana Tymchyshyn. Probabilistic model checking of complex biological pathways. *Theor. Comput. Sci.*, 391(3):239–257, 2008.