

Translating Stochastic CLS into Maude (ONGOING WORK)

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Introduction

The Calculus of Looping Sequences (CLS) is a formalism for the description of biological systems

- Stochastic CLS is the stochastic extension of CLS
- A simulator based on Stochastic CLS has been developed

Our original aim was to apply model checking to Stochastic CLS models

We wanted to use Probabilistic Maude (PMaude) as a model checker

- unfortunately, the model checking module of PMaude seems not to be available...

Consequently,

- we have chosen Real-Time Maude (Maude with a notion of time)
- we have adapted Gillespie's stochastic simulation algorithm in order to be used in Real-Time Maude
- we have used Real-Time Maude analysis tool to verify properties on the results of a number of simulations (statistical model checking)

Outline of the talk

1 Introduction

2 The Calculus of Looping Sequences (CLS)

- Definition of CLS
- The *lac* operon in CLS
- Stochastic CLS

3 (Statistical) model checking of Stochastic CLS models

- Choosing a model checker
- Translation of Stochastic CLS into Real-Time Maude
- Analysis examples

The Calculus of Looping Sequences (CLS)

We assume an alphabet \mathcal{E} . **Terms** T and **Sequences** S of CLS are given by the following grammar:

$$\begin{aligned} T &::= S \mid (S)^L \mid T \mid T \\ S &::= \epsilon \mid a \mid S \cdot S \end{aligned}$$

where a is a generic element of \mathcal{E} , and ϵ is the empty sequence.

The operators are:

$S \cdot S$: Sequencing

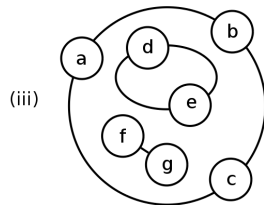
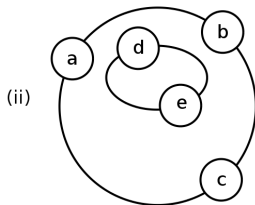
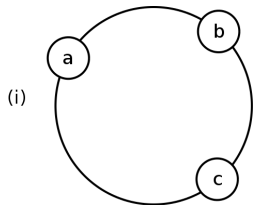
$(S)^L$: Looping (S is closed and it can rotate)

$T_1 \mid T_2$: Containment (T_1 contains T_2)

$T \mid T$: Parallel composition (juxtaposition)

Actually, looping and containment form a single binary operator $(S)^L \mid T$.

Examples of Terms



$$(i) \quad (a \cdot b \cdot c)^L \rfloor \epsilon$$

$$(ii) \quad (a \cdot b \cdot c)^L \rfloor (d \cdot e)^L \rfloor \epsilon$$

$$(iii) \quad (a \cdot b \cdot c)^L \rfloor (f \cdot g \mid (d \cdot e)^L) \rfloor \epsilon$$

Structural Congruence

The **Structural Congruence** relations \equiv_S and \equiv_T are the least congruence relations on sequences and on terms, respectively, satisfying the following rules:

$$S_1 \cdot (S_2 \cdot S_3) \equiv_S (S_1 \cdot S_2) \cdot S_3 \quad S \cdot \epsilon \equiv_S \epsilon \cdot S \equiv_S S$$

$$T_1 \mid T_2 \equiv_T T_2 \mid T_1 \quad T_1 \mid (T_2 \mid T_3) \equiv_T (T_1 \mid T_2) \mid T_3$$

$$T \mid \epsilon \equiv_T T \quad (\epsilon)^L \rfloor \epsilon \equiv_T \epsilon \quad (S_1 \cdot S_2)^L \rfloor T \equiv_T (S_2 \cdot S_1)^L \rfloor T$$

We write \equiv for \equiv_T .

CLS Patterns

Let us consider variables of three kinds:

- term variables (X, Y, Z, \dots)
- sequence variables ($\tilde{x}, \tilde{y}, \tilde{z}, \dots$)
- element variables (x, y, z, \dots)

Patterns P and **Sequence Patterns** SP of CLS extend CLS terms and sequences with variables:

$$\begin{aligned} P & ::= SP \mid (SP)^L \mid P \mid P \mid X \\ SP & ::= \epsilon \mid a \mid SP \cdot SP \mid x \mid \tilde{x} \end{aligned}$$

where a is a generic element of \mathcal{E} , ϵ is the empty sequence, and x, \tilde{x} and X are generic element, sequence and term variables

The structural congruence relation \equiv extends trivially to patterns

Rewrite Rules

$P\sigma$ denotes the term obtained by replacing any variable in T with the corresponding term, sequence or element.

Σ is the set of all possible instantiations σ

A **Rewrite Rule** is a pair (P, P') , denoted $P \mapsto P'$, where:

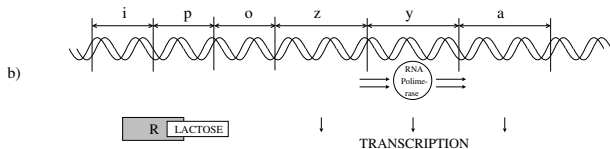
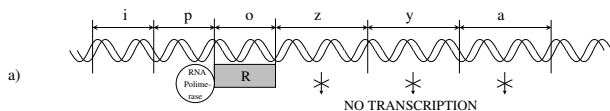
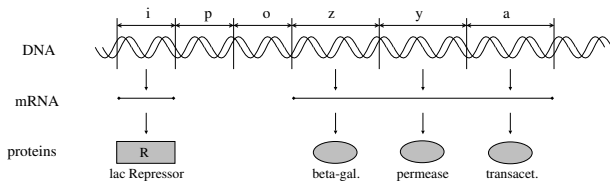
- P, P' are patterns
- variables in P' are a subset of those in P

A rule $P \mapsto P'$ can be applied to all terms $P\sigma$.

Example: $a \cdot x \cdot a \mapsto b \cdot x \cdot b$

- can be applied to $a \cdot c \cdot a$ (producing $b \cdot c \cdot b$)
- cannot be applied to $a \cdot c \cdot c \cdot a$

CLS modeling examples: the *lac* operon (1)



CLS modeling examples: the *lac* operon (2)

$$Ecoli ::= (m)^L \mid (lacI \cdot lacP \cdot lacO \cdot lacZ \cdot lacY \cdot lacA \mid polym)$$

Rules for DNA transcription/translation:

$$lacI \cdot \tilde{x} \mapsto lacI' \cdot \tilde{x} \mid repr \quad (R1)$$
$$polym \mid \tilde{x} \cdot lacP \cdot \tilde{y} \mapsto \tilde{x} \cdot PP \cdot \tilde{y} \quad (R2)$$
$$\tilde{x} \cdot PP \cdot lacO \cdot \tilde{y} \mapsto \tilde{x} \cdot lacP \cdot PO \cdot \tilde{y} \quad (R3)$$
$$\tilde{x} \cdot PO \cdot lacZ \cdot \tilde{y} \mapsto \tilde{x} \cdot lacO \cdot PZ \cdot \tilde{y} \quad (R4)$$
$$\tilde{x} \cdot PZ \cdot lacY \cdot \tilde{y} \mapsto \tilde{x} \cdot lacZ \cdot PY \cdot \tilde{y} \mid betagal \quad (R5)$$
$$\tilde{x} \cdot PY \cdot lacA \mapsto \tilde{x} \cdot lacY \cdot PA \mid perm \quad (R6)$$
$$\tilde{x} \cdot PA \mapsto \tilde{x} \cdot lacA \mid transac \mid polym \quad (R7)$$

CLS modeling examples: the *lac* operon (3)

$$Ecoli ::= (m)^L \rfloor (lacI \cdot lacP \cdot lacO \cdot lacZ \cdot lacY \cdot lacA \mid polym)$$

Rules to describe the binding of the lac Repressor to gene o, and what happens when lactose is present in the environment of the bacterium:

$$repr \mid \tilde{x} \cdot lacO \cdot \tilde{y} \mapsto \tilde{x} \cdot RO \cdot \tilde{y} \quad (R8)$$

$$LACT \mid (m \cdot \tilde{x})^L \rfloor X \mapsto (m \cdot \tilde{x})^L \rfloor (X \mid LACT) \quad (R9)$$

$$\tilde{x} \cdot RO \cdot \tilde{y} \mid LACT \mapsto \tilde{x} \cdot lacO \cdot \tilde{y} \mid RLACT \quad (R10)$$

$$(\tilde{x})^L \rfloor (perm \mid X) \mapsto (perm \cdot \tilde{x})^L \rfloor X \quad (R11)$$

$$LACT \mid (perm \cdot \tilde{x})^L \rfloor X \mapsto (perm \cdot \tilde{x})^L \rfloor (LACT \mid X) \quad (R12)$$

$$betagal \mid LACT \mapsto betagal \mid GLU \mid GAL \quad (R13)$$

CLS modeling examples: the *lac* operon (4)

$$Ecoli ::= (m)^L \rfloor (lacI \cdot lacP \cdot lacO \cdot lacZ \cdot lacY \cdot lacA \mid polym)$$

Example:

$$Ecoli \mid LACT \mid LACT$$
$$\rightarrow^* (m)^L \rfloor (lacI' \cdot lacP \cdot lacO \cdot lacZ \cdot lacY \cdot lacA \mid polym \mid repr) \mid LACT \mid LACT$$
$$\rightarrow^* (m)^L \rfloor (lacI' \cdot lacP \cdot RO \cdot lacZ \cdot lacY \cdot lacA \mid polym) \mid LACT \mid LACT$$
$$\rightarrow^* (m)^L \rfloor (lacI' \cdot lacP \cdot lacO \cdot lacZ \cdot lacY \cdot lacA \mid polym \mid RLACT) \mid LACT$$
$$\rightarrow^* (perm \cdot m)^L \rfloor (lacI' - A \mid betagal \mid transac \mid polym \mid RLACT) \mid LACT$$
$$\rightarrow^* (perm \cdot m)^L \rfloor (lacI' - A \mid betagal \mid transac \mid polym \mid RLACT \mid GLU \mid GAL)$$

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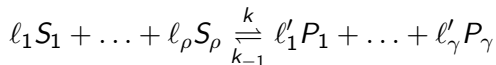
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Background: the kinetics of chemical reactions

Usual notation for chemical reactions:



where:

- S_i, P_i are molecules (reactants)
- ℓ_i, ℓ'_i are stoichiometric coefficients
- k, k_{-1} are the kinetic constants

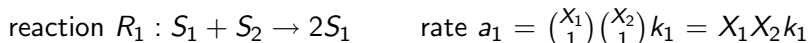
The kinetics is described by the *law of mass action*:

$$\frac{d[P_i]}{dt} = \underbrace{\ell'_i k [S_1]^{\ell_1} \dots [S_\rho]^{\ell_\rho}}_{\text{reaction rate}} - \underbrace{\ell'_i k_{-1} [P_1]^{\ell'_1} \dots [P_\gamma]^{\ell'_\gamma}}_{\text{reaction rate}}$$

Background: Gillespie's simulation algorithm

- represents a chemical solution as a multiset of molecules
- computes the reaction rate a_μ by multiplying the kinetic constant by the number of possible combinations of reactants

Example: chemical solution with X_1 molecules S_1 and X_2 molecules S_2



Given a set of reactions $\{R_1, \dots, R_M\}$ and a current time t

- The time $t + \tau$ at which the next reaction will occur is randomly chosen with τ exponentially distributed with parameter $\sum_{\nu=1}^M a_\nu$;
- The reaction R_μ that has to occur at time $t + \tau$ is randomly chosen with probability $\frac{a_\mu}{\sum_{\nu=1}^M a_\nu}$.

At each step t is incremented by τ and the chemical solution is updated.

Stochastic CLS

Stochastic CLS incorporates Gillespie's stochastic framework into the semantics of CLS

- Rewrite rules are enriched with a kinetic constant

What is a reactant combination in Stochastic CLS?

- A *reactant combination* is an occurrence (up to \equiv) of a left hand side of a rewrite rule

Two definitions of the semantics of Stochastic CLS exist

- The first computes the number of reactant combinations by adding unique labels to the alphabet symbols in the term
- The second computes the number of reactant combinations compositionally

The *occ* function

For the sake of the translation into Maude we have defined a recursive function $occ(T, T')$ that gives the number of combinations of reactants T in the term T' .

$$occ(T, \epsilon) = 0$$

$$occ(S, E \cdot S_1 | T) = \begin{cases} 1 + occ(S, S_1) + occ(S, T) & \text{if } S \text{ is a prefix of } E \cdot S_1 \\ occ(S, S_1) + occ(S, T) & \text{otherwise} \end{cases}$$

$$occ(S, (S_1)^L | T) | T_1 = occ'(S, S_1) + occ(S, T) + occ(S, T_1)$$

$$occ((S)^L | T_1, (S_1)^L | T | T_2) = \begin{cases} 1 + occ((S)^L | T_1, T_2) & \text{if } S \equiv S_1 \text{ and } T_1 \equiv T \\ occ((S)^L | T_1, T) + occ((S)^L | T_1, T_2) & \text{otherwise} \end{cases}$$

$$occ(S | T, S | T_1) = \begin{cases} \frac{(1 + exactocc(S, T_1)) exactocc(T, T_1)}{1 + exactocc(S, T)} & \text{if } exactocc(T, T_1) > 0 \\ occ(S | T, T_1) & \text{otherwise} \end{cases}$$

$$occ((S)^L | T | T_1, (S_1)^L | T_2 | T_3) = \begin{cases} \frac{(1 + exactocc((S)^L | T, T_3)) exactocc(T_1, T_3)}{1 + exactocc((S)^L | T, T_1)} & \text{if } S \equiv S_1, T \equiv T_2 \\ & \text{and } exactocc(T_1, T_3) > 0 \\ occ((S)^L | T | T_1, T_2) + occ((S)^L | T | T_1, T_3) & \text{otherwise} \end{cases}$$

$$\vdots$$
$$\vdots$$
$$\vdots$$
$$\vdots$$

The occ function: examples

$$\text{occ}(a | b , a | a | b | b | b) = 6$$

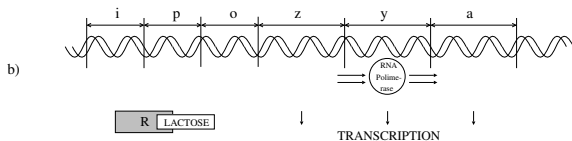
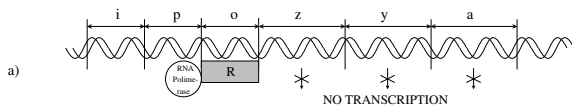
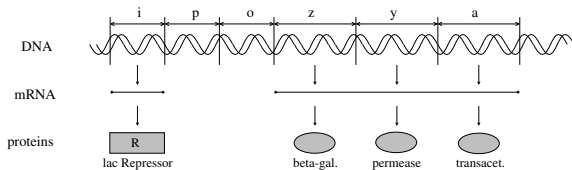
$$\text{occ}((m)^L \rfloor a , (m)^L \rfloor a | (m)^L \rfloor (m)^L \rfloor a) = 2$$

$$\text{occ}(a | b , (m)^L \rfloor (a | b) | (m)^L \rfloor (a | b | a)) = 3$$

$$\text{occ}(a \cdot b , a \cdot b \cdot c \cdot a \cdot b) = 2$$

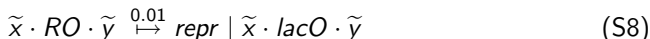
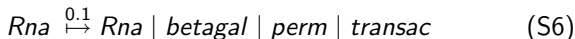
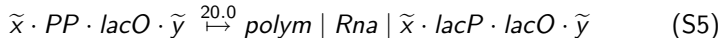
$$\text{occ}(a \cdot b , (b \cdot a \cdot b \cdot a)^L \rfloor a \cdot b \cdot c) = 3$$

A Stochastic CLS model of the *lac* operon (1)



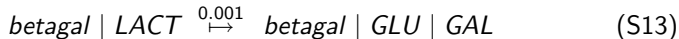
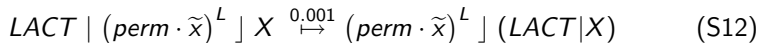
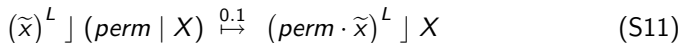
A Stochastic CLS model of the *lac* operon (2)

Transcription of DNA, binding of lac Repressor to gene *o*, and interaction between lactose and lac Repressor:

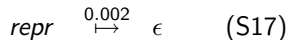
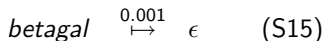


A Stochastic CLS model of the *lac* operon (3)

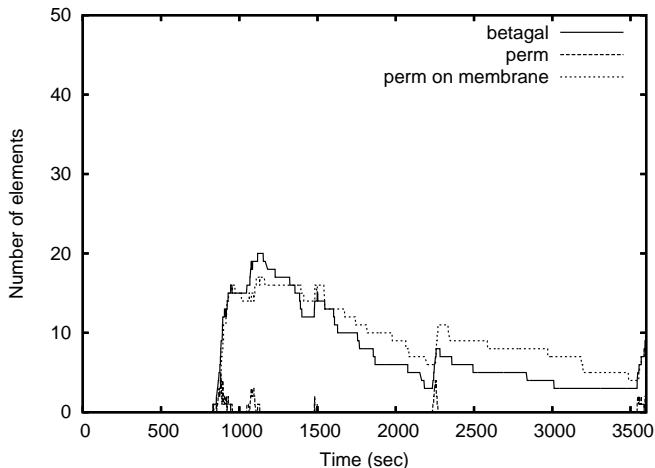
The behaviour of the three enzymes for lactose degradation:



Degradation of all the proteins and mRNA involved in the process:

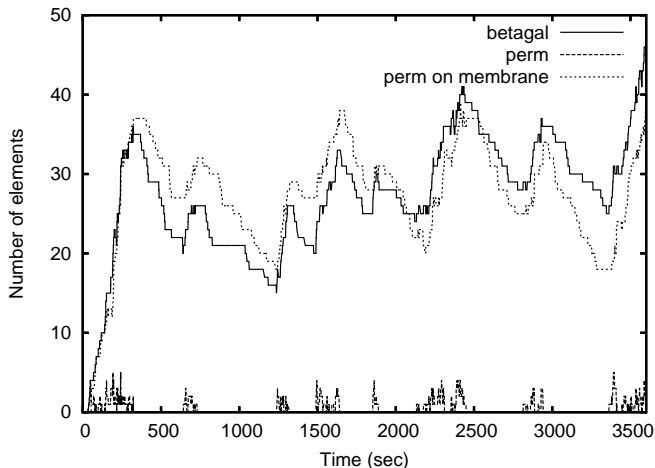


Simulation results (1)



Production of enzymes in the absence of lactose
 $(m)^L \rfloor (lacl - A \mid 30 \times polym \mid 100 \times repr)$

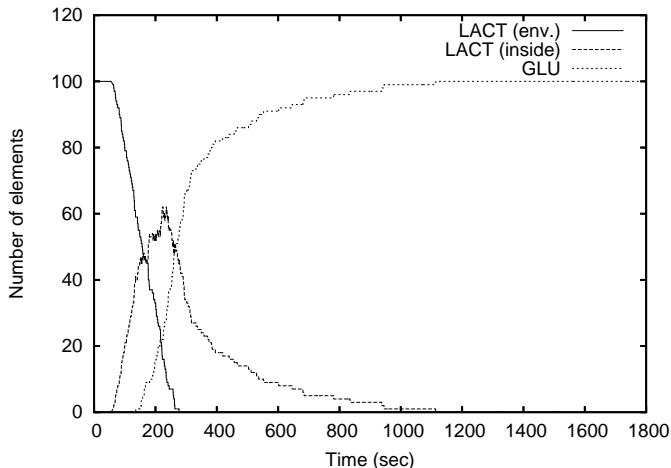
Simulation results (2)



Production of enzymes in the presence of lactose

$100 \times LACT \mid (m)^L \mid (lacI - A \mid 30 \times polym \mid 100 \times repr)$

Simulation results (3)



Degradation of lactose into glucose

$100 \times LACT \mid (m)^L \mid (lact - A \mid 30 \times polym \mid 100 \times repr)$

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A model checker for Stochastic CLS

As candidate model checkers we have considered:

- PRISM
- Murphi
- PMaude

All of them are probabilistic/stochastic model checkers

PMaude is the most suitable

- It uses a language based on rewrite rules (rewrite logic) that eases the translation of Stochastic CLS rules

Unfortunately, the model checking module of PMaude seems not to be available

- a possible alternative: Real-Time Maude

Real-Time Maude

Maude is a specification language equipped with efficient analysis tools, which supports three modelling paradigms:

- algebraic style (via equations)
- rewrite logic (via rewrite rules)
- object oriented (via classes and messages)

Real-Time Maude extends Maude with a notion of time

- rewrite rule applications might consume (a fixed amount of) time

Real-Time Maude has two kinds of rules

- instantaneous rules:

`cr1 [l] : t => t' if cond`

- tick rules:

`cr1 [l] : t => t' in time τ if cond`

Translation of Stochastic CLS into Real-Time Maude

Real-Time Maude is not stochastic

- we will include Gillespie's simulation algorithm (slightly changed) in the translation of Stochastic CLS models
- it will be used to generate single executions of the model
- Real-Time Maude analysis tools will be applied to the simulation results

This is *statistical model checking*

- we loose exhaustivity (properties are checked on a number of runs)
- huge systems could be handled
- may allow *property driven* simulations

A slight modification to Gillespie's algorithm

Given reactions $\{R_1, \dots, R_M\}$ and initial molecular population X_1, \dots, X_M :

Step 0 Initialize current time $t = 0$ and stepped time $t_step = 0$

Step 1 Compute reaction propensities a_1, \dots, a_n and $\sum_{\nu=1}^M a_\nu$

Step 2 Choose reaction μ and reaction time τ

Step 3 If $t + \tau \geq t_step$ then $t_step = t_step + \Delta t$

Step 4 Execute reaction μ and set $t = t + \tau$

Step 5 If $t \geq total_time$ then conclude, else return to **Step 1**

Simulations are still exact

- our modification doesn't change states and choices of the simulation
- simulation results are sampled every Δt time units

(Modified) Gillespie's algorithm in Real-Time Maude

Every step of the simulation algorithm is translated into an instantaneous rule, apart from **Step 3** (the increase of the stepped time)

```
(set tick def 1/100 .)
```

```
cr1 [increase] :  
  < step:3 , t:R1 , tau:R2 , t_step:R3 , delta_t:R4 >  
=>  
  < step:4 , t_step:R3+R4 >  
in time R3  
if (R1+R2) >= R3
```

```
cr1 [no_increase] :  
  < step:3 , t:R1 , tau:R2 , t_step:R3 , delta_t:R4 >  
=>  
  < step:4 >  
in time R3  
if (R1+R2) < R3
```

Translation of Stochastic CLS into Real-Time Maude

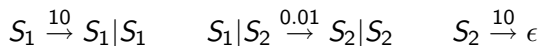
$$\begin{aligned} T & ::= S \mid (S)^L \mid T \mid T \\ S & ::= \epsilon \mid a \mid S \cdot S \end{aligned}$$

```
(omod CLS is
  pr NAT
  sorts Elem Seq Term Loop
  subsorts Elem < Seq < Term

  op empty : -> Seq [ctor]
  op ... : Seq Seq -> Seq
    [assoc gather (E e) id: empty ctor]
  op '{_}'_ : Elem Nat -> Term
  op '['_']LContains '['_'] : Seq Term -> Term
    [prec 41 gather (& &) ctor]
  op '|_' : Term Term -> Term
    [assoc comm prec 45 gather (E e) id: empty ctor]
endom)
```

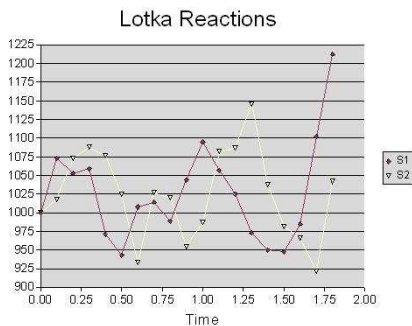
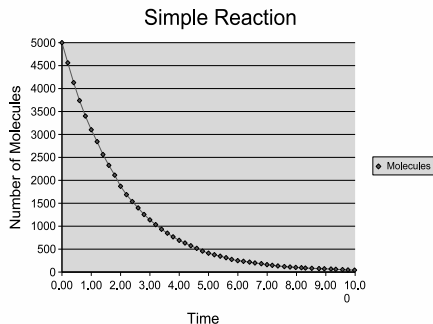
Translation of Stochastic CLS into Real-Time Maude

Lotka reactions as Stochastic CLS rules



```
r1 [ S1 ] :  
  < 0 : CLSTerm | term : (T | S1), mu : 1, step : 4 >  
=>  
  < 0 : CLSTerm | term : (T | S1 | S1), step : 5 >  
  
r1 [ S2 ] :  
  < 0 : CLSTerm | term : (T | S1 | S2), mu : 2, step : 4 >  
=>  
  < 0 : CLSTerm | term : (T | S2 | S2), step : 5 >  
  
r1 [ S3 ] :  
  < 0 : CLSTerm | term : (T | S2), mu : 3, step : 4 >  
=>  
  < 0 : CLSTerm | term : T, step : 5 >
```


Analysis example: stochastic simulation



• Simple reaction: $S_1 \xrightarrow{0.5} S_2$

• Lotka reactions: $S_1 \xrightarrow{10} S_1|S_1$ $S_1|S_2 \xrightarrow{0.01} S_2|S_2$ $S_2 \xrightarrow{10} \epsilon$

Analysis example: statistical model checking

Initialisation of 100 stochastic simulations

```
rl [ initialise1 ] :  
  < step : 0 >  
=>  
  < seed : random(1), step : 1 >  
:  
:  
:  
rl [ initialise100 ] :  
  < step : 0 >  
=>  
  < seed : random(100), step : 1 >
```

Analysis example: statistical model checking

By using the `tsearch` command we can check all possible behaviours

Starting with $4 \times S_1$ and $4 \times S_2$ we search 10 states where S_2 is absent

```
(tsearch [10] INIT({S1}4 | {S2}4) =>* {< 0:0id : CLSTerm | term
: T:Term > C:Configuration} such that occ(S2,T:Term) = 0 in time
<= 1/10 .)
```

Solution 1

```
C:Configuration -->
```

```
< term: {S1}5, finaltime: 7.8293318117206676e-2 >
```

```
:
```

Solution 10

```
C:Configuration -->
```

```
< term: {S1}8, finaltime: 5.6307762323583766e-2 >
```

Analysis example: statistical model checking

The `find earliest` searches for the earliest time when a given state is reached

Starting with $4 \times S_1$ and $4 \times S_2$ we search the earliest time when S_2 disappear

```
(find earliest INIT({S1}4 | {S2}4) =>* {< CLSTerm | term:T:Term >  
C:Configuration} such that occ(S2,T:Term) == 0 .)
```

Result:

```
{< term: {S1}8, finaltime: 5.6307762323583766e-2 > in time 3/50}
```

Analysis example: statistical model checking

Verification of properties expressed as LTL formulas. Some state formulas:

`vanished(T)` indicates that term `T` has vanished from the system,

`IsLessThan(T,T')` indicates that the occurrences of term `T` are less than the occurrences of `T'`.

Starting with $4 \times S_1$ and $4 \times S_2$ we prove

- that S_2 will eventually disappear (i.e. $\diamond \text{vanished}(S_2)$)
- that the amount of S_2 will eventually become less than the amount of S_1 (i.e. $\diamond \text{IsLessThan}(S_2, S_1)$)

```
(mc INIT({S1}4 | {S2}4) |=t <> vanished(S2) in time<=1 .)
```

Result Bool : true

```
(mc INIT({S1} 4 | {S2} 4) |=t <> IsLessThan(S2,S1) in time<=1 .)
```

Result Bool : true

Conclusions

We have presented preliminary ideas for the application of (statistical) model checking to Stochastic CLS models

Many things to do:

- Prove the correctness of the translation into Real-Time Maude w.r.t. Stochastic CLS semantics
- Improve the efficiency of the computation of the *occ* function
- Model and analyse some more significant case study (e.g. the Lactose operon)