## **SBML Model Report**

# Model name: "Tyson2003\_Perfect\_Adaption"



May 5, 2016

## 1 General Overview

This is a document in SBML Level 2 Version 4 format. This model was created by Lukas Endler<sup>1</sup> at February tenth 2011 at 2:38 a.m. and last time modified at April sixth 2014 at 8:08 p.m. Table 1 provides an overview of the quantities of all components of this model.

Table 1: Number of components in this model, which are described in the following sections.

Element	Quantity	Element	Quantity
compartment types	0	compartments	1
species types	0	species	3
events	0	constraints	0
reactions	4	function definitions	0
global parameters	5	unit definitions	6
rules	1	initial assignments	1

#### **Model Notes**

This is an SBML implementation the model of the perfect adaptor (figure 1d) described in the article:

Sniffers, buzzers, toggles and blinkers: dynamics of regulatory and signaling pathways in the cell.

Tyson JJ, Chen KC, Novak B. <u>Curr Opin Cell Biol.</u> 2003 Apr;15(2):221-31. PubmedID:12648679; DOI:10.1016/S0955-0674(03)00017-6;

<sup>&</sup>lt;sup>1</sup>EMBL-EBI, lukas@ebi.ac.uk

#### Abstract:

The physiological responses of cells to external and internal stimuli are governed by genes and proteins interacting in complex networks whose dynamical properties are impossible to understand by intuitive reasoning alone. Recent advances by theoretical biologists have demonstrated that molecular regulatory networks can be accurately modeled in mathematical terms. These models shed light on the design principles of biological control systems and make predictions that have been verified experimentally.

Originally created by libAntimony v1.4 (using libSBML 3.4.1)

This model originates from BioModels Database: A Database of Annotated Published Models (http://www.ebi.ac.uk/biomodels/). It is copyright (c) 2005-2011 The BioModels.net Team. For more information see the terms of use.

To cite BioModels Database, please use: Li C, Donizelli M, Rodriguez N, Dharuri H, Endler L, Chelliah V, Li L, He E, Henry A, Stefan MI, Snoep JL, Hucka M, Le Novre N, Laibe C (2010) BioModels Database: An enhanced, curated and annotated resource for published quantitative kinetic models. BMC Syst Biol., 4:92.

## 2 Unit Definitions

This is an overview of nine unit definitions of which three are predefined by SBML and not mentioned in the model.

#### 2.1 Unit time

Name s

**Definition** s

#### 2.2 Unit substance

Name mole

**Definition** mol

### 2.3 Unit per\_s

Name per\_s

**Definition**  $s^{-1}$ 

## 2.4 Unit M\_per\_s

Name M\_per\_s

**Definition**  $mol \cdot s^{-1} \cdot l^{-1}$ 

### 2.5 Unit M

Name M

**Definition**  $mol \cdot l^{-1}$ 

## 2.6 Unit per\_M\_per\_s

Name per\_M\_per\_s

**Definition**  $\text{mol}^{-1} \cdot 1 \cdot \text{s}^{-1}$ 

### 2.7 Unit volume

**Notes** Litre is the predefined SBML unit for volume.

**Definition** 1

#### 2.8 Unit area

**Notes** Square metre is the predefined SBML unit for area since SBML Level 2 Version 1.

**Definition** m<sup>2</sup>

### 2.9 Unit length

**Notes** Metre is the predefined SBML unit for length since SBML Level 2 Version 1.

**Definition** m

## 3 Compartment

This model contains one compartment.

Table 2: Properties of all compartments.

Id	Name	SBO	Spatial Dimensions	Size	Unit	Constant	Outside
env		0000270	3	1	litre	<b>✓</b>	

## 3.1 Compartment env

This is a three dimensional compartment with a constant size of one litre.

**SBO:0000270** enzymatic rate law for competitive inhibition of irreversible unireactant enzymes by exclusive inhibitors

# 4 Species

This model contains three species. The boundary condition of one of these species is set to true so that this species' amount cannot be changed by any reaction. Section 9 provides further details and the derived rates of change of each species.

Table 3: Properties of each species.

Id	Name	Compartment	Derived Unit	Constant	Boundary Condi- tion
R		env	$\operatorname{mol} \cdot \operatorname{l}^{-1}$	$\Box$	
X		env	$\text{mol} \cdot l^{-1}$		
S		env	$\text{mol} \cdot l^{-1}$		$\square$

## **5 Parameters**

This model contains five global parameters.

Table 4: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
k1		0000035	2.0	$s^{-1}$	$\checkmark$
k2		0000036	2.0	$\text{mol}^{-1} \cdot 1 \cdot \text{s}^{-1}$	
k3		0000035	1.0	$s^{-1}$	
k4		0000356	1.0	$s^{-1}$	$\square$
tau		0000002	4.0	S	

## 6 Initialassignment

This is an overview of one initial assignment.

## **6.1 Initialassignment** R

Derived unit  $mol \cdot l^{-1}$ 

Math  $\frac{k1 \cdot k4}{k2 \cdot k3}$ 

## 7 Rule

This is an overview of one rule.

### **7.1 Rule S**

Rule S is an assignment rule for species S:

$$S = 1 \cdot \left\lfloor \frac{\text{time}}{\text{tau}} \right\rfloor \tag{1}$$

6

## 8 Reactions

This model contains four reactions. All reactions are listed in the following table and are subsequently described in detail. If a reaction is affected by a modifier, the identifier of this species is written above the reaction arrow.

Table 5: Overview of all reactions

N⁰	Id	Name	Reaction Equation	SBO
1	r1		$\emptyset \xrightarrow{S} R$	0000176
2	r2		$R \xrightarrow{X} \emptyset$	0000179
3 4	r3 r4		$ \emptyset \xrightarrow{S} X  X \longrightarrow \emptyset $	0000176 0000179

### 8.1 Reaction r1

This is an irreversible reaction of no reactant forming one product influenced by one modifier.

SBO:0000176 biochemical reaction

## **Reaction equation**

$$\emptyset \xrightarrow{S} R$$
 (2)

**Modifier** 

Table 6: Properties of each modifier.

Id	Name	SBO
S		0000461

#### **Product**

Table 7: Properties of each product.

Id	Name	SBO
R		

## **Kinetic Law**

SBO:0000049 mass action rate law for first order irreversible reactions, continuous scheme

Derived unit  $s^{-1} \cdot mol$ 

$$v_1 = \text{vol}(\text{env}) \cdot \text{k1} \cdot [S] \tag{3}$$

### 8.2 Reaction r2

This is an irreversible reaction of one reactant forming no product influenced by one modifier.

SBO:0000179 degradation

## **Reaction equation**

$$R \xrightarrow{X} \emptyset \tag{4}$$

Reactant

Table 8: Properties of each reactant.

Id	Name	SBO
R		

### **Modifier**

Table 9: Properties of each modifier.

Id	Name	SBO
X		0000461

### **Kinetic Law**

**SBO:0000054** mass action rate law for second order irreversible reactions, two reactants, continuous scheme

Derived unit  $s^{-1} \cdot mol$ 

$$v_2 = \text{vol}(\text{env}) \cdot \text{k2} \cdot [\text{R}] \cdot [\text{X}] \tag{5}$$

## 8.3 Reaction r3

This is an irreversible reaction of no reactant forming one product influenced by one modifier.

SBO:0000176 biochemical reaction

## **Reaction equation**

$$\emptyset \xrightarrow{S} X \tag{6}$$

### **Modifier**

Table 10: Properties of each modifier.

Id	Name	SBO
S		0000461

#### **Product**

Table 11: Properties of each product.

### **Kinetic Law**

SBO:0000049 mass action rate law for first order irreversible reactions, continuous scheme

**Derived unit**  $s^{-1} \cdot mol$ 

$$v_3 = \text{vol}(\text{env}) \cdot \text{k3} \cdot [S] \tag{7}$$

### 8.4 Reaction r4

This is an irreversible reaction of one reactant forming no product.

SBO:0000179 degradation

### **Reaction equation**

$$X \longrightarrow \emptyset$$
 (8)

#### Reactant

Table 12: Properties of each reactant.

Id	Name	SBO
Х		

#### **Kinetic Law**

SBO:0000049 mass action rate law for first order irreversible reactions, continuous scheme

**Derived unit**  $s^{-1} \cdot mol$ 

$$v_4 = \text{vol}(\text{env}) \cdot \text{k4} \cdot [X] \tag{9}$$

## 9 Derived Rate Equations

When interpreted as an ordinary differential equation framework, this model implies the following set of equations for the rates of change of each species.

## 9.1 Species R

SBO:0000252 polypeptide chain

#### Initial assignment R

This species takes part in two reactions (as a reactant in r2 and as a product in r1).

$$\frac{\mathrm{d}}{\mathrm{d}t}\mathbf{R} = v_1 - v_2 \tag{10}$$

#### 9.2 Species X

SBO:0000252 polypeptide chain

Initial concentration  $0 \text{ mol} \cdot l^{-1}$ 

This species takes part in three reactions (as a reactant in r4 and as a product in r3 and as a modifier in r2).

$$\frac{\mathrm{d}}{\mathrm{d}t}\mathbf{X} = v_3 - v_4 \tag{11}$$

## 9.3 Species S

SBO:0000285 material entity of unspecified nature

#### Involved in rule S

This species takes part in two reactions (as a modifier in r1, r3). Not these but one rule determines the species' quantity because this species is on the boundary of the reaction system.

## A Glossary of Systems Biology Ontology Terms

- **SBO:0000002** quantitative systems description parameter: A numerical value that defines certain characteristics of systems or system functions. It may be part of a calculation, but its value is not determined by the form of the equation itself, and may be arbitrarily assigned
- **SBO:0000035 forward unimolecular rate constant, continuous case:** Numerical parameter that quantifies the forward velocity of a chemical reaction involving only one reactant. This parameter encompasses all the contributions to the velocity except the quantity of the reactant. It is to be used in a reaction modelled using a continuous framework
- **SBO:0000036 forward bimolecular rate constant, continuous case:** Numerical parameter that quantifies the forward velocity of a chemical reaction involving two reactants. This parameter encompasses all the contributions to the velocity except the quantity of the reactants. It is to be used in a reaction modelled using a continuous framework

- SBO:0000049 mass action rate law for first order irreversible reactions, continuous scheme:
  - Reaction scheme where the products are created from the reactants and the change of a product quantity is proportional to the product of reactant activities. The reaction scheme does not include any reverse process that creates the reactants from the products. The change of a product quantity is proportional to the quantity of one reactant. It is to be used in a reaction modelled using a continuous framework.
- **SBO:0000054** mass action rate law for second order irreversible reactions, two reactants, continuous scheme: Reaction scheme where the products are created from the reactants and the change of a product quantity is proportional to the product of reactant activities. The reaction scheme does not include any reverse process that creates the reactants from the products. The change of a product quantity is proportional to the product of two reactant quantities. It is to be used in a reaction modelled using a continuous framework.
- **SBO:0000176 biochemical reaction:** An event involving one or more chemical entities that modifies the electrochemical structure of at least one of the participants.
- **SBO:0000179 degradation:** Complete disappearance of a physical entity
- **SBO:0000252** polypeptide chain: Naturally occurring macromolecule formed by the repetition of amino-acid residues linked by peptidic bonds. A polypeptide chain is synthesized by the ribosome. CHEBI:1654
- **SBO:0000270** enzymatic rate law for competitive inhibition of irreversible unireactant enzymes by exclusive inhibitors: Inhibition of a unireactant enzyme by inhibitors that bind to the free enzyme on the same binding site than the substrate. The enzymes do not catalyse the reactions in both directions.
- **SBO:0000285 material entity of unspecified nature:** Material entity whose nature is unknown or irrelevant
- **SBO:0000356 decay constant:** Kinetic constant characterising a mono-exponential decay. It is the inverse of the mean lifetime of the continuant being decayed. Its unit is "per tim".
- **SBO:0000461 essential activator:** A substance that is absolutely required for occurrence and stimulation of a reaction

SML2ATEX was developed by Andreas Dräger<sup>a</sup>, Hannes Planatscher<sup>a</sup>, Dieudonné M Wouamba<sup>a</sup>, Adrian Schröder<sup>a</sup>, Michael Hucka<sup>b</sup>, Lukas Endler<sup>c</sup>, Martin Golebiewski<sup>d</sup> and Andreas Zell<sup>a</sup>. Please see http://www.ra.cs.uni-tuebingen.de/software/SBML2LaTeX for more information.

<sup>&</sup>lt;sup>a</sup>Center for Bioinformatics Tübingen (ZBIT), Germany

<sup>&</sup>lt;sup>b</sup>California Institute of Technology, Beckman Institute BNMC, Pasadena, United States

<sup>&</sup>lt;sup>c</sup>European Bioinformatics Institute, Wellcome Trust Genome Campus, Hinxton, United Kingdom

<sup>&</sup>lt;sup>d</sup>EML Research gGmbH, Heidelberg, Germany