

SBML Model Report

Model name: “Mandlik2015 - Tristable genetic circuit of Leishmania”



May 6, 2016

1 General Overview

This is a document in SBML Level 2 Version 1 format. This model was created by the following two authors: Shailza Singh¹ and Vijayalakshmi Chelliah² at October 26th 2015 at 12:57 a. m. and last time modified at February twelveth 2016 at 3:21 p. m. Table 1 gives 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	35
events	0	constraints	0
reactions	14	function definitions	0
global parameters	34	unit definitions	0
rules	14	initial assignments	0

2 Unit Definitions

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

¹National Centre for Cell Science, Pune, singhs@nccs.res.in

²EMBL-EBI, viji@ebi.ac.uk

2.1 Unit substance

Notes Mole is the predefined SBML unit for substance.

Definition mol

2.2 Unit volume

Notes Litre is the predefined SBML unit for volume.

Definition l

2.3 Unit area

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

Definition m²

2.4 Unit length

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

Definition m

2.5 Unit time

Notes Second is the predefined SBML unit for time.

Definition s

3 Compartment

This model contains one compartment.

Table 2: Properties of all compartments.

Id	Name	SBO	Spatial Dimensions	Size	Unit	Constant	Outside
DefaultCompartment	DefaultCompartment		3	1	litre	<input checked="" type="checkbox"/>	

3.1 Compartment DefaultCompartment

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

Name DefaultCompartment

4 Species

This model contains 35 species. The boundary condition of 14 of these species is set to true so that these species' amount cannot be changed by any reaction. Section 8 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 Condition
AUR1	AUR1	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
IPTG	IPTG	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
LACI	LACI	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
LAMDAR	LAMDAR	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
SLS1	SLS1	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
SLS4	SLS4	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
TETR	TETR	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
as1	as1	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
ia1_ActiveTF	ia1_ActiveTF	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
ia1_InactiveTF	ia1_InactiveTF	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
ope1	ope1	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
p1	p1	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
p2	p2	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
p3	p3	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
pAUR1	pAUR1	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
pSLS1	pSLS1	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
pSLS4	pSLS4	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
proAUR1	proAUR1	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
proLACI	proLACI	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
proLAMDAR	proLAMDAR	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
proSLS1	proSLS1	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>

Id	Name	Compartment	Derived Unit	Constant	Boundary Condi- tion
proSLS4	proSLS4	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
proTETR	proTETR	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
rs1	rs1	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
rs2	rs2	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
rs3	rs3	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
rs4	rs4	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
rs5	rs5	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
rs6	rs6	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
ter1	ter1	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
ter2	ter2	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
ter3	ter3	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
ter4	ter4	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
ter5	ter5	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
ter6	ter6	DefaultCompartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>

5 Parameters

This model contains 34 global parameters.

Table 4: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
tr3_Kd	tr3_Kd		1.000		✓
tr3_h	tr3_h		2.000		✓
tr4_Kd	tr4_Kd		1.000		✓
tr4_h	tr4_h		2.000		✓
pSLS4- _strength	pSLS4_strength		2.570		✓
tr5_Kd	tr5_Kd		1.000		✓
tr5_h	tr5_h		2.000		✓
tr6_Kd	tr6_Kd		1.000		✓
tr6_h	tr6_h		2.000		✓
p1_strength	p1_strength		2.500		✓
p2_strength	p2_strength		2.400		✓
proLACI- _degradation- _rate	proLACI- _degradation_rate		0.070		✓
p3_strength	p3_strength		2.500		✓
proSLS1- _degradation- _rate	proSLS1- _degradation_rate		0.070		✓
proSLS4- _degradation- _rate	proSLS4- _degradation_rate		0.040		✓
proLAMDAR- _degradation- _rate	proLAMDAR- _degradation_rate		0.042		✓
proAUR1- _degradation- _rate	proAUR1- _degradation_rate		0.100		✓
ia1_Kd	ia1_Kd		0.001		✓
ia1- _repression- _Kd	ia1_repression_Kd		1.000		✓
ia1- _repression_h	ia1_repression_h		2.000		✓
pAUR1- _strength	pAUR1_strength		2.014		✓

Id	Name	SBO	Value	Unit	Constant
proTETR- _degradation- _rate	proTETR- _degradation_rate		0.080		✓
ta1_Kd	ta1_Kd		1.000		✓
ta1_h	ta1_h		2.000		✓
pp2- _translation- _rate	pp2_translation- _rate		1.000		✓
pp3- _translation- _rate	pp3_translation- _rate		1.000		✓
pp4- _translation- _rate	pp4_translation- _rate		1.000		✓
pp5- _translation- _rate	pp5_translation- _rate		1.000		✓
pp6- _translation- _rate	pp6_translation- _rate		1.000		✓
tr1_Kd	tr1_Kd		1.000		✓
tr1_h	tr1_h		2.000		✓
pSLS1- _strength	pSLS1_strength		2.500		✓
tr2_Kd	tr2_Kd		1.000		✓
tr2_h	tr2_h		2.000		✓

6 Rules

This is an overview of 14 rules.

6.1 Rule rs6

Rule rs6 is an assignment rule for species rs6:

$$rs6 = \frac{1}{1 + \left(\frac{[proTETR]}{tr4_Kd} \right)^{tr4_h}} \quad (1)$$

6.2 Rule rs3

Rule rs3 is an assignment rule for species rs3:

$$rs3 = \frac{1}{1 + \left(\frac{[proLACI]}{tr6.Kd} \right)^{tr6.h}} \quad (2)$$

6.3 Rule rs4

Rule rs4 is an assignment rule for species rs4:

$$rs4 = \frac{1}{1 + \left(\frac{[proLAMDAR]}{tr5.Kd} \right)^{tr5.h}} \quad (3)$$

6.4 Rule rs5

Rule rs5 is an assignment rule for species rs5:

$$rs5 = \frac{1}{1 + \left(\frac{[proLACI]}{tr3.Kd} \right)^{tr3.h}} \quad (4)$$

6.5 Rule LAMDAR

Rule LAMDAR is an assignment rule for species LAMDAR:

$$LAMDAR = p1_strength \cdot [rs6] \quad (5)$$

6.6 Rule LACI

Rule LACI is an assignment rule for species LACI:

$$LACI = p2_strength \cdot [rs4] \quad (6)$$

6.7 Rule TETR

Rule TETR is an assignment rule for species TETR:

$$TETR = p3_strength \cdot [rs5] \quad (7)$$

6.8 Rule rs1

Rule rs1 is an assignment rule for species rs1:

$$rs1 = \frac{1}{1 + \left(\frac{[proSLS1]}{tr1.Kd} \right)^{tr1.h}} \quad (8)$$

6.9 Rule rs2

Rule rs2 is an assignment rule for species rs2:

$$rs2 = \frac{1}{1 + \left(\frac{[proAUR1]}{tr2_Kd} \right)^{tr2_h}} \quad (9)$$

6.10 Rule ope1

Rule ope1 is an assignment rule for species ope1:

$$ope1 = \frac{1}{1 + \left(\frac{[ia1_ActiveTF]}{ia1_repression_Kd} \right)^{ia1_repression_h}} \quad (10)$$

6.11 Rule as1

Rule as1 is an assignment rule for species as1:

$$as1 = \frac{1 + \left(\frac{[proSLS4]}{ta1_Kd} \right)^{ta1_h} - 1}{1 + \left(\frac{[proSLS4]}{ta1_Kd} \right)^{ta1_h}} \quad (11)$$

6.12 Rule SLS1

Rule SLS1 is an assignment rule for species SLS1:

$$SLS1 = pSLS1_strength \cdot [as1] \cdot [ope1] \cdot [rs3] \quad (12)$$

6.13 Rule SLS4

Rule SLS4 is an assignment rule for species SLS4:

$$SLS4 = pSLS4_strength \cdot [rs2] \quad (13)$$

6.14 Rule AUR1

Rule AUR1 is an assignment rule for species AUR1:

$$AUR1 = pAUR1_strength \cdot [rs1] \quad (14)$$

7 Reactions

This model contains 14 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

Nº	Id	Name	Reaction Equation	SBO
1	pp1.v1	pp1.v1	$\emptyset \longrightarrow \text{proSLS1}$	
2	pp1.v2	pp1.v2	$\text{proSLS1} \xrightarrow{\text{proSLS1}} \emptyset$	
3	pp5.v1	pp5.v1	$\emptyset \xrightarrow{\text{LAMDAR}, \text{LAMDAR}} \text{proLAMDAR}$	
4	pp5.v2	pp5.v2	$\text{proLAMDAR} \xrightarrow{\text{proLAMDAR}} \emptyset$	
5	ia1.bind- _forward	ia1_bind_forward	$\text{IPTG} + \text{ia1_ActiveTF} \xrightarrow{\text{IPTG}, \text{ia1_ActiveTF}} \text{ia1_InactiveTF}$	
6	ia1.bind- _reverse	ia1_bind_reverse	$\text{ia1_InactiveTF} \xrightarrow{\text{ia1_InactiveTF}} \text{IPTG}$ ia1_ActiveTF	+
7	pp2.v1	pp2.v1	$\emptyset \xrightarrow{\text{AUR1}, \text{AUR1}} \text{proAUR1}$	
8	pp2.v2	pp2.v2	$\text{proAUR1} \xrightarrow{\text{proAUR1}} \emptyset$	
9	pp6.v1	pp6.v1	$\emptyset \xrightarrow{\text{TETR}, \text{TETR}} \text{proTETR}$	
10	pp6.v2	pp6.v2	$\text{proTETR} \xrightarrow{\text{proTETR}} \emptyset$	
11	pp3.v1	pp3.v1	$\emptyset \xrightarrow{\text{SLS4}, \text{SLS4}} \text{proSLS4}$	
12	pp3.v2	pp3.v2	$\text{proSLS4} \xrightarrow{\text{proSLS4}} \emptyset$	
13	pp4.v1	pp4.v1	$\emptyset \xrightarrow{\text{LACI}, \text{LACI}} \text{proLACI}$	
14	pp4.v2	pp4.v2	$\text{proLACI} \xrightarrow{\text{proLACI}} \emptyset$	

7.1 Reaction pp1_v1

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

Name pp1_v1

Reaction equation



Product

Table 6: Properties of each product.

Id	Name	SBO
proSLS1	proSLS1	

Kinetic Law

Derived unit not available

$$v_1 = 1 \quad (16)$$

7.2 Reaction pp1_v2

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

Name pp1_v2

Reaction equation



Reactant

Table 7: Properties of each reactant.

Id	Name	SBO
proSLS1	proSLS1	

Modifier

Table 8: Properties of each modifier.

Id	Name	SBO
proSLS1	proSLS1	

Kinetic Law

Derived unit contains undeclared units

$$v_2 = \text{proSLS1_degradation_rate} \cdot [\text{proSLS1}] \quad (18)$$

7.3 Reaction pp5_v1

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

Name pp5_v1

Reaction equation



Modifiers

Table 9: Properties of each modifier.

Id	Name	SBO
LAMDAR	LAMDAR	
LAMDAR	LAMDAR	

Product

Table 10: Properties of each product.

Id	Name	SBO
proLAMDAR	proLAMDAR	

Kinetic Law

Derived unit contains undeclared units

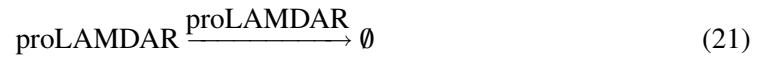
$$v_3 = \text{pp5_translation_rate} \cdot \text{p1_strength} \cdot [\text{LAMDAR}] \quad (20)$$

7.4 Reaction pp5_v2

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

Name pp5_v2

Reaction equation



Reactant

Table 11: Properties of each reactant.

Id	Name	SBO
proLAMDAR	proLAMDAR	

Modifier

Table 12: Properties of each modifier.

Id	Name	SBO
proLAMDAR	proLAMDAR	

Kinetic Law

Derived unit contains undeclared units

$$v_4 = \text{proLAMDAR_degradation_rate} \cdot [\text{proLAMDAR}] \quad (22)$$

7.5 Reaction ia1_bind_forward

This is an irreversible reaction of two reactants forming one product influenced by two modifiers.

Name ia1_bind_forward

Reaction equation



Reactants

Table 13: Properties of each reactant.

Id	Name	SBO
IPTG	IPTG	
ia1_ActiveTF	ia1_ActiveTF	

Modifiers

Table 14: Properties of each modifier.

Id	Name	SBO
IPTG	IPTG	
ia1_ActiveTF	ia1_ActiveTF	

Product

Table 15: Properties of each product.

Id	Name	SBO
ia1_InactiveTF	ia1_InactiveTF	

Kinetic Law

Derived unit $\text{mol}^2 \cdot \text{l}^{-2}$

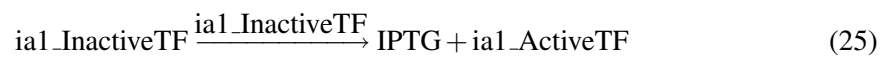
$$v_5 = [\text{IPTG}] \cdot [\text{ia1_ActiveTF}] \quad (24)$$

7.6 Reaction ia1_bind_reverse

This is an irreversible reaction of one reactant forming two products influenced by one modifier.

Name ia1_bind_reverse

Reaction equation



Reactant

Table 16: Properties of each reactant.

Id	Name	SBO
ia1_InactiveTF	ia1_InactiveTF	

Modifier

Table 17: Properties of each modifier.

Id	Name	SBO
ia1_InactiveTF	ia1_InactiveTF	

Products

Table 18: Properties of each product.

Id	Name	SBO
IPTG	IPTG	
ia1_ActiveTF	ia1_ActiveTF	

Kinetic Law

Derived unit contains undeclared units

$$v_6 = \text{ia1_Kd} \cdot [\text{ia1_InactiveTF}] \quad (26)$$

7.7 Reaction pp2_v1

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

Name pp2_v1

Reaction equation



Modifiers

Table 19: Properties of each modifier.

Id	Name	SBO
AUR1	AUR1	
AUR1	AUR1	

Product

Table 20: Properties of each product.

Id	Name	SBO
proAUR1	proAUR1	

Kinetic Law

Derived unit contains undeclared units

$$v_7 = \text{pp2_translation_rate} \cdot \text{pAUR1_strength} \cdot [\text{AUR1}] \quad (28)$$

7.8 Reaction pp2_v2

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

Name pp2_v2

Reaction equation



Reactant

Table 21: Properties of each reactant.

Id	Name	SBO
proAUR1	proAUR1	

Modifier

Table 22: Properties of each modifier.

Id	Name	SBO
proAUR1	proAUR1	

Kinetic Law

Derived unit contains undeclared units

$$v_8 = \text{proAUR1_degradation_rate} \cdot [\text{proAUR1}] \quad (30)$$

7.9 Reaction pp6_v1

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

Name pp6_v1

Reaction equation



Modifiers

Table 23: Properties of each modifier.

Id	Name	SBO
TETR	TETR	
TETR	TETR	

Product

Table 24: Properties of each product.

Id	Name	SBO
proTETR	proTETR	

Kinetic Law

Derived unit contains undeclared units

$$v_9 = \text{pp6_translation_rate} \cdot \text{p3_strength} \cdot [\text{TETR}] \quad (32)$$

7.10 Reaction pp6_v2

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

Name pp6_v2

Reaction equation



Reactant

Table 25: Properties of each reactant.

Id	Name	SBO
proTETR	proTETR	

Modifier

Table 26: Properties of each modifier.

Id	Name	SBO
proTETR	proTETR	

Kinetic Law

Derived unit contains undeclared units

$$v_{10} = \text{proTETR_degradation_rate} \cdot [\text{proTETR}] \quad (34)$$

7.11 Reaction pp3_v1

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

Name pp3_v1

Reaction equation



Modifiers

Table 27: Properties of each modifier.

Id	Name	SBO
SLS4	SLS4	
SLS4	SLS4	

Product

Table 28: Properties of each product.

Id	Name	SBO
proSLS4	proSLS4	

Kinetic Law

Derived unit contains undeclared units

$$v_{11} = \text{pp3_translation_rate} \cdot \text{pSLS4_strength} \cdot [\text{SLS4}] \quad (36)$$

7.12 Reaction pp3_v2

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

Name pp3_v2

Reaction equation



Reactant

Table 29: Properties of each reactant.

Id	Name	SBO
proSLS4	proSLS4	

Modifier

Table 30: Properties of each modifier.

Id	Name	SBO
proSLS4	proSLS4	

Kinetic Law

Derived unit contains undeclared units

$$v_{12} = \text{proSLS4_degradation_rate} \cdot [\text{proSLS4}] \quad (38)$$

7.13 Reaction pp4_v1

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

Name pp4_v1

Reaction equation



Modifiers

Table 31: Properties of each modifier.

Id	Name	SBO
LACI	LACI	
LACI	LACI	

Product

Table 32: Properties of each product.

Id	Name	SBO
proLACI	proLACI	

Kinetic Law

Derived unit contains undeclared units

$$v_{13} = \text{pp4_translation_rate} \cdot \text{p2_strength} \cdot [\text{LACI}] \quad (40)$$

7.14 Reaction pp4_v2

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

Name pp4_v2

Reaction equation



Reactant

Table 33: Properties of each reactant.

Id	Name	SBO
proLACI	proLACI	

Modifier

Table 34: Properties of each modifier.

Id	Name	SBO
proLACI	proLACI	

Kinetic Law

Derived unit contains undeclared units

$$v_{14} = \text{proLACI.degradation_rate} \cdot [\text{proLACI}] \quad (42)$$

8 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.

Identifiers for kinetic laws highlighted in gray cannot be verified to evaluate to units of SBML substance per time. As a result, some SBML interpreters may not be able to verify the consistency of the units on quantities in the model. Please check if

- parameters without an unit definition are involved or
- volume correction is necessary because the `hasOnlySubstanceUnits` flag may be set to `false` and `spacialDimensions` > 0 for certain species.

8.1 Species AUR1

Name AUR1

Initial concentration $10 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule AUR1

This species takes part in two reactions (as a modifier in [pp2.v1](#), [pp2.v1](#)). Not these but one rule determines the species' quantity because this species is on the boundary of the reaction system.

8.2 Species IPTG

Name IPTG

Initial concentration $10000 \text{ mol} \cdot \text{l}^{-1}$

This species takes part in three reactions (as a reactant in [ia1.bind.forward](#) and as a product in [ia1.bind.reverse](#) and as a modifier in [ia1.bind.forward](#)).

$$\frac{d}{dt} \text{IPTG} = v_6 - v_5 \quad (43)$$

8.3 Species LACI

Name LACI

Initial concentration $10 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule LACI

This species takes part in two reactions (as a modifier in [pp4.v1](#), [pp4.v1](#)). Not these but one rule determines the species' quantity because this species is on the boundary of the reaction system.

8.4 Species LAMDAR

Name LAMDAR

Initial concentration $10 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule LAMDAR

This species takes part in two reactions (as a modifier in [pp5.v1](#), [pp5.v1](#)). Not these but one rule determines the species' quantity because this species is on the boundary of the reaction system.

8.5 Species SLS1

Name SLS1

Initial concentration $10 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule SLS1

One rule determines the species' quantity.

8.6 Species SLS4

Name SLS4

Initial concentration $10 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule SLS4

This species takes part in two reactions (as a modifier in [pp3_v1](#), [pp3_v1](#)). Not these but one rule determines the species' quantity because this species is on the boundary of the reaction system.

8.7 Species TETR

Name TETR

Initial concentration $10 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule TETR

This species takes part in two reactions (as a modifier in [pp6_v1](#), [pp6_v1](#)). Not these but one rule determines the species' quantity because this species is on the boundary of the reaction system.

8.8 Species as1

Name as1

Initial concentration $10 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule as1

One rule determines the species' quantity.

8.9 Species ia1_ActiveTF

Name ia1_ActiveTF

Initial concentration $100 \text{ mol} \cdot \text{l}^{-1}$

This species takes part in three reactions (as a reactant in [ia1_bind_forward](#) and as a product in [ia1_bind_reverse](#) and as a modifier in [ia1_bind_forward](#)).

$$\frac{d}{dt} \text{ia1_ActiveTF} = v_6 - v_5 \quad (44)$$

8.10 Species `ia1_InactiveTF`

Name `ia1_InactiveTF`

Initial concentration $100 \text{ mol} \cdot \text{l}^{-1}$

This species takes part in three reactions (as a reactant in `ia1_bind_reverse` and as a product in `ia1_bind_forward` and as a modifier in `ia1_bind_reverse`).

$$\frac{d}{dt} \text{ia1_InactiveTF} = v_5 - v_6 \quad (45)$$

8.11 Species `ope1`

Name `ope1`

Initial concentration $5 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule `ope1`

One rule determines the species' quantity.

8.12 Species `p1`

Name `p1`

Initial concentration $10 \text{ mol} \cdot \text{l}^{-1}$

This species does not take part in any reactions. Its quantity does hence not change over time:

$$\frac{d}{dt} p1 = 0 \quad (46)$$

8.13 Species `p2`

Name `p2`

Initial concentration $10 \text{ mol} \cdot \text{l}^{-1}$

This species does not take part in any reactions. Its quantity does hence not change over time:

$$\frac{d}{dt} p2 = 0 \quad (47)$$

8.14 Species `p3`

Name `p3`

Initial concentration $10 \text{ mol} \cdot \text{l}^{-1}$

This species does not take part in any reactions. Its quantity does hence not change over time:

$$\frac{d}{dt} p3 = 0 \quad (48)$$

8.15 Species p_{AUR1}

Name p_{AUR1}

Initial concentration $10 \text{ mol} \cdot \text{l}^{-1}$

This species does not take part in any reactions. Its quantity does hence not change over time:

$$\frac{d}{dt}p_{\text{AUR1}} = 0 \quad (49)$$

8.16 Species p_{SLS1}

Name p_{SLS1}

Initial concentration $10 \text{ mol} \cdot \text{l}^{-1}$

This species does not take part in any reactions. Its quantity does hence not change over time:

$$\frac{d}{dt}p_{\text{SLS1}} = 0 \quad (50)$$

8.17 Species p_{SLS4}

Name p_{SLS4}

Initial concentration $10 \text{ mol} \cdot \text{l}^{-1}$

This species does not take part in any reactions. Its quantity does hence not change over time:

$$\frac{d}{dt}p_{\text{SLS4}} = 0 \quad (51)$$

8.18 Species pro_{AUR1}

Name pro_{AUR1}

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

This species takes part in three reactions (as a reactant in [pp2_v2](#) and as a product in [pp2_v1](#) and as a modifier in [pp2_v2](#)).

$$\frac{d}{dt}pro_{\text{AUR1}} = v_7 - v_8 \quad (52)$$

8.19 Species `proLACI`

Name `proLACI`

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

This species takes part in three reactions (as a reactant in `pp4_v2` and as a product in `pp4_v1` and as a modifier in `pp4_v2`).

$$\frac{d}{dt}\text{proLACI} = v_{13} - v_{14} \quad (53)$$

8.20 Species `proLAMДАР`

Name `proLAMДАР`

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

This species takes part in three reactions (as a reactant in `pp5_v2` and as a product in `pp5_v1` and as a modifier in `pp5_v2`).

$$\frac{d}{dt}\text{proLAMДАР} = v_3 - v_4 \quad (54)$$

8.21 Species `proSLS1`

Name `proSLS1`

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

This species takes part in three reactions (as a reactant in `pp1_v2` and as a product in `pp1_v1` and as a modifier in `pp1_v2`).

$$\frac{d}{dt}\text{proSLS1} = v_1 - v_2 \quad (55)$$

8.22 Species `proSLS4`

Name `proSLS4`

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

This species takes part in three reactions (as a reactant in `pp3_v2` and as a product in `pp3_v1` and as a modifier in `pp3_v2`).

$$\frac{d}{dt}\text{proSLS4} = v_{11} - v_{12} \quad (56)$$

8.23 Species `proTETR`

Name `proTETR`

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

This species takes part in three reactions (as a reactant in `pp6_v2` and as a product in `pp6_v1` and as a modifier in `pp6_v2`).

$$\frac{d}{dt} \text{proTETR} = v_9 - v_{10} \quad (57)$$

8.24 Species `rs1`

Name `rs1`

Initial concentration $10 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule `rs1`

One rule determines the species' quantity.

8.25 Species `rs2`

Name `rs2`

Initial concentration $10 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule `rs2`

One rule determines the species' quantity.

8.26 Species `rs3`

Name `rs3`

Initial concentration $10 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule `rs3`

One rule determines the species' quantity.

8.27 Species `rs4`

Name `rs4`

Initial concentration $10 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule `rs4`

One rule determines the species' quantity.

8.28 Species `rs5`

Name `rs5`

Initial concentration $10 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule `rs5`

One rule determines the species' quantity.

8.29 Species `rs6`

Name `rs6`

Initial concentration $10 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule `rs6`

One rule determines the species' quantity.

8.30 Species `ter1`

Name `ter1`

Initial concentration $10 \text{ mol} \cdot \text{l}^{-1}$

This species does not take part in any reactions. Its quantity does hence not change over time:

$$\frac{d}{dt} \text{ter1} = 0 \quad (58)$$

8.31 Species `ter2`

Name `ter2`

Initial concentration $10 \text{ mol} \cdot \text{l}^{-1}$

This species does not take part in any reactions. Its quantity does hence not change over time:

$$\frac{d}{dt} \text{ter2} = 0 \quad (59)$$

8.32 Species `ter3`

Name `ter3`

Initial concentration $10 \text{ mol} \cdot \text{l}^{-1}$

This species does not take part in any reactions. Its quantity does hence not change over time:

$$\frac{d}{dt} \text{ter3} = 0 \quad (60)$$

8.33 Species `ter4`

Name `ter4`

Initial concentration $10 \text{ mol} \cdot \text{l}^{-1}$

This species does not take part in any reactions. Its quantity does hence not change over time:

$$\frac{d}{dt}\text{ter4} = 0 \quad (61)$$

8.34 Species `ter5`

Name `ter5`

Initial concentration $10 \text{ mol} \cdot \text{l}^{-1}$

This species does not take part in any reactions. Its quantity does hence not change over time:

$$\frac{d}{dt}\text{ter5} = 0 \quad (62)$$

8.35 Species `ter6`

Name `ter6`

Initial concentration $10 \text{ mol} \cdot \text{l}^{-1}$

This species does not take part in any reactions. Its quantity does hence not change over time:

$$\frac{d}{dt}\text{ter6} = 0 \quad (63)$$

SBML2^{AT}EX was developed by Andreas Dräger^a, Hannes Planatscher^a, Dieudonné M Wouamba^a, Adrian Schröder^a, Michael Hucka^b, Lukas Endler^c, Martin Golebiewski^d and Andreas Zell^a. Please see <http://www.ra.cs.uni-tuebingen.de/software/SBML2LaTeX> for more information.

^aCenter for Bioinformatics Tübingen (ZBIT), Germany

^bCalifornia Institute of Technology, Beckman Institute BNMC, Pasadena, United States

^cEuropean Bioinformatics Institute, Wellcome Trust Genome Campus, Hinxton, United Kingdom

^dEML Research gGmbH, Heidelberg, Germany