

SBML Model Report

Model name: “Obeyesekere1999_CellCycle”



May 6, 2016

1 General Overview

This is a document in SBML Level 2 Version 1 format. This model was created by Harish Dharuri¹ at March 16th 2008 at 7:31 p. m. and last time modified at May 27th 2014 at 10:24 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	7
events	0	constraints	0
reactions	10	function definitions	0
global parameters	21	unit definitions	0
rules	3	initial assignments	0

Model Notes

The model reproduces the time profiles of the different species depicted in Fig 3a of the paper. Model successfully reproduced using MathSBML.

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2 Unit Definitions

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

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
cell_1	cell		3	1	litre	<input checked="" type="checkbox"/>	

3.1 Compartment `cell_1`

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

Name `cell`

4 Species

This model contains seven species. 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
D_1	D	cell_1	$\text{mol} \cdot \text{l}^{-1}$	\square	\square
E_1	E	cell_1	$\text{mol} \cdot \text{l}^{-1}$	\square	\square
RS_1	RS	cell_1	$\text{mol} \cdot \text{l}^{-1}$	\square	\square
R_1	R	cell_1	$\text{mol} \cdot \text{l}^{-1}$	\square	\square
X_1	X	cell_1	$\text{mol} \cdot \text{l}^{-1}$	\square	\square
E2F_1	E2F	cell_1	$\text{mol} \cdot \text{l}^{-1}$	\square	\square
RP_1	RP	cell_1	$\text{mol} \cdot \text{l}^{-1}$	\square	\square

5 Parameters

This model contains 21 global parameters.

Table 4: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
aD_1	aD		0.400		<input checked="" type="checkbox"/>
k_1	k		0.050		<input checked="" type="checkbox"/>
GF_1	GF		6.300		<input checked="" type="checkbox"/>
dD_1	dD		0.400		<input checked="" type="checkbox"/>
aE_1	aE		0.160		<input checked="" type="checkbox"/>
af_1	af		0.900		<input checked="" type="checkbox"/>
theta_1	theta		1.500		<input checked="" type="checkbox"/>
dE_1	dE		0.200		<input checked="" type="checkbox"/>
pX_1	pX		0.480		<input checked="" type="checkbox"/>
RT_1	RT		2.500		<input checked="" type="checkbox"/>
pS_1	pS		0.600		<input checked="" type="checkbox"/>
pD_1	pD		0.480		<input checked="" type="checkbox"/>
qD_1	qD		0.600		<input checked="" type="checkbox"/>
pE_1	pE		0.096		<input checked="" type="checkbox"/>
qE_1	qE		0.600		<input checked="" type="checkbox"/>
aX_1	aX		0.080		<input checked="" type="checkbox"/>
f_1	f		0.200		<input checked="" type="checkbox"/>
g_1	g		0.528		<input checked="" type="checkbox"/>
dX_1	dX		1.040		<input checked="" type="checkbox"/>
qX_1	qX		0.800		<input checked="" type="checkbox"/>
unpho_RB			0.000		<input type="checkbox"/>

6 Rules

This is an overview of three rules.

6.1 Rule E2F_1

Rule E2F_1 is an assignment rule for species E2F_1:

$$E2F_1 = \text{theta}_1 - [RS_1] \quad (1)$$

6.2 Rule RP_1

Rule RP_1 is an assignment rule for species RP_1:

$$RP_1 = RT_1 - [RS_1] - [R_1] \quad (2)$$

6.3 Rule `unpho_RB`

Rule `unpho_RB` is an assignment rule for parameter `unpho_RB`:

$$\text{unpho_RB} = [\text{R_1}] + [\text{RS_1}] \quad (3)$$

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

7 Reactions

This model contains ten 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	_1	cycD/CDK4 synthesis	$\emptyset \longrightarrow D_1$	
2	cyclinCDK4degradation_1	cycD/CDK4 degradation	$D_1 \xrightarrow{D_1, E_1} \emptyset$	
3	cyclin_1	cycE/CDK2 synthesis	$\emptyset \xrightarrow{E2F_1} E_1$	
4	cycECDK2degradation_1	cycE/CDK2 degradation	$E_1 \xrightarrow{E_1, X_1} \emptyset$	
5	pRBpdephosphorylation_1	pRBp dephosphorilation	$\emptyset \xrightarrow{RP_1, X_1} R_1$	
6	pRBE2Fcomplexassociation_1	pRB/E2F-complex association	$R_1 \xrightarrow{RS_1, R_1, E2F_1} RS_1$	
7	pRBE2Fcomplexdeassociation_1	pRB/E2F-complex deassociation via cycD/CDK4	$RS_1 \xrightarrow{RS_1, D_1} \emptyset$	
8	_7	pRB/E2F complex deassociation via cycE/CDK2	$RS_1 \xrightarrow{RS_1, E_1} \emptyset$	
9	cycleprogression_1	cycle progression	$\emptyset \xrightarrow{E_1, E2F_1, X_1} X_1$	
10	cyclebreak_1	cycle break	$X_1 \xrightarrow{X_1} \emptyset$	

7.1 Reaction _1

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

Name cycD/CDK4 synthesis

Reaction equation



Product

Table 6: Properties of each product.

Id	Name	SBO
D_1	D	

Kinetic Law

Derived unit not available

$$v_1 = aD_1 \cdot \frac{k_1 \cdot GF_1}{1 + k_1 \cdot GF_1} \quad (5)$$

7.2 Reaction cyclinCDK4degradation_1

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

Name cycD/CDK4 degradation

Reaction equation



Reactant

Table 7: Properties of each reactant.

Id	Name	SBO
D_1	D	

Modifiers

Table 8: Properties of each modifier.

Id	Name	SBO
D_1	D	
E_1	E	

Kinetic Law

Derived unit contains undeclared units

$$v_2 = dD_1 \cdot [E_1] \cdot [D_1] \quad (7)$$

7.3 Reaction `cyclin_1`

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

Name `cycE/CDK2 synthesis`

Reaction equation



Modifier

Table 9: Properties of each modifier.

Id	Name	SBO
E2F_1	E2F	

Product

Table 10: Properties of each product.

Id	Name	SBO
E_1	E	

Kinetic Law

Derived unit contains undeclared units

$$v_3 = aE_1 \cdot (1 + af_1 \cdot [E2F_1]) \quad (9)$$

7.4 Reaction [cycECDK2degradation_1](#)

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

Name cycE/CDK2 degradation

Reaction equation



Reactant

Table 11: Properties of each reactant.

Id	Name	SBO
E_1	E	

Modifiers

Table 12: Properties of each modifier.

Id	Name	SBO
E_1	E	
X_1	X	

Kinetic Law

Derived unit contains undeclared units

$$v_4 = dE_1 \cdot [X_1] \cdot [E_1] \quad (11)$$

7.5 Reaction [pRBpdephosphorylation_1](#)

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

Name pRB-p dephosphorilation

Reaction equation



Modifiers

Table 13: Properties of each modifier.

Id	Name	SBO
RP_1	RP	
X_1	X	

Product

Table 14: Properties of each product.

Id	Name	SBO
R_1	R	

Kinetic Law

Derived unit contains undeclared units

$$v_5 = \frac{p_{X_1} \cdot [RP_1] \cdot [X_1]}{q_{X_1} + [RP_1] + [X_1]} \quad (13)$$

7.6 Reaction `pRB/E2Fcomplexassociation_1`

This is an irreversible reaction of one reactant forming one product influenced by three modifiers.

Name pRB/E2F complex association

Reaction equation



Reactant

Table 15: Properties of each reactant.

Id	Name	SBO
R_1	R	

Modifiers

Table 16: Properties of each modifier.

Id	Name	SBO
RS_1	RS	
R_1	R	
E2F_1	E2F	

Product

Table 17: Properties of each product.

Id	Name	SBO
RS_1	RS	

Kinetic Law

Derived unit contains undeclared units

$$v_6 = pS_1 \cdot [E2F_1] \cdot [R_1] \quad (15)$$

7.7 Reaction pRBE2FcomplexdeassociationviacycDCDK4_1

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

Name pRB/E2F complex deassociation via cycD/CDK4

Reaction equation



Reactant

Table 18: Properties of each reactant.

Id	Name	SBO
RS_1	RS	

Modifiers

Table 19: Properties of each modifier.

Id	Name	SBO
RS_1	RS	
D_1	D	

Kinetic Law

Derived unit contains undeclared units

$$v_7 = \frac{pD_1 \cdot [RS_1] \cdot [D_1]}{qD_1 + [RS_1] + [D_1]} \quad (17)$$

7.8 Reaction _7

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

Name pRB/E2F complex deassociation via cycE/CDK2

Reaction equation



Reactant

Table 20: Properties of each reactant.

Id	Name	SBO
RS_1	RS	

Modifiers

Table 21: Properties of each modifier.

Id	Name	SBO
RS_1	RS	
E_1	E	

Kinetic Law

Derived unit contains undeclared units

$$v_8 = \frac{pE_{-1} \cdot [RS_{-1}] \cdot [E_{-1}]}{qE_{-1} + [RS_{-1}] + [E_{-1}]} \quad (19)$$

7.9 Reaction `cycleprogression_1`

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

Name cycle progression

Reaction equation



Modifiers

Table 22: Properties of each modifier.

Id	Name	SBO
E ₋₁	E	
E2F ₋₁	E2F	
X ₋₁	X	

Product

Table 23: Properties of each product.

Id	Name	SBO
X ₋₁	X	

Kinetic Law

Derived unit contains undeclared units

$$v_9 = aX_{-1} \cdot [E_{-1}] + f_{-1} \cdot [E2F_{-1}] + g_{-1} \cdot [X_{-1}]^2 \cdot [E_{-1}] \quad (21)$$

7.10 Reaction `cyclebreak_1`

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

Name cycle break

Reaction equation



Reactant

Table 24: Properties of each reactant.

Id	Name	SBO
X_1	X	

Modifier

Table 25: Properties of each modifier.

Id	Name	SBO
X_1	X	

Kinetic Law

Derived unit contains undeclared units

$$v_{10} = dX_1 \cdot [X_1] \quad (23)$$

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 D_1

Name D

Initial amount 0.1 mol

This species takes part in four reactions (as a reactant in [cyclinCDK4degradation_1](#) and as a product in [_1](#) and as a modifier in [cyclinCDK4degradation_1](#), [prBE2FcomplexdeassociationviacycDCDK4-1](#)).

$$\frac{d}{dt}D_1 = v_1 - v_2 \quad (24)$$

8.2 Species E_1

Name E

Initial amount 0.6 mol

This species takes part in six reactions (as a reactant in [cycECDK2degradation_1](#) and as a product in [cyclin_1](#) and as a modifier in [cyclinCDK4degradation_1](#), [cycECDK2degradation_1](#), [_7](#), [cycleprogression_1](#)).

$$\frac{d}{dt}E_1 = v_3 - v_4 \quad (25)$$

8.3 Species RS_1

Name RS

Initial amount 1 mol

This species takes part in six reactions (as a reactant in [pRBE2FcomplexdeassociationviacycDCDK4_1](#), [_7](#) and as a product in [pRBE2Fcomplexassociation_1](#) and as a modifier in [pRBE2Fcomplexassociation_1](#), [pRBE2FcomplexdeassociationviacycDCDK4_1](#), [_7](#)).

$$\frac{d}{dt}RS_1 = v_6 - v_7 - v_8 \quad (26)$$

8.4 Species R_1

Name R

Initial amount 0.5 mol

This species takes part in three reactions (as a reactant in [pRBE2Fcomplexassociation_1](#) and as a product in [pRBpdephosphorylation_1](#) and as a modifier in [pRBE2Fcomplexassociation_1](#)).

$$\frac{d}{dt}R_1 = v_5 - v_6 \quad (27)$$

8.5 Species X_1

Name X

Initial amount 0.7 mol

This species takes part in six reactions (as a reactant in [cyclebreak_1](#) and as a product in [cycleprogression_1](#) and as a modifier in [cycECDK2degradation_1](#), [pRBpdephosphorylation_1](#), [cycleprogression_1](#), [cyclebreak_1](#)).

$$\frac{d}{dt}X_1 = v_9 - v_{10} \quad (28)$$

8.6 Species E2F_1

Name E2F

Involved in rule E2F_1

This species takes part in three reactions (as a modifier in [cyclin_1](#), [pRBE2Fcomplexassociation_1](#), [cycleprogression_1](#)) and is also involved in one rule which determines this species' quantity.

8.7 Species RP_1

Name RP

Involved in rule RP_1

This species takes part in one reaction (as a modifier in [pRBpdephosphorylation_1](#)) and is also involved in one rule which determines this species' quantity.

SBML²TeX 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.

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