

SBML Model Report

Model name: “Bai2003_G1phaseRegulation”



May 5, 2016

1 General Overview

This is a document in SBML Level 2 Version 4 format. This model was created by the following two authors: Vijayalakshmi Chelliah¹ and Mandri Obeyesekere² at March fifth 2010 at 4:22 p. m. and last time modified at February 14th 2014 at 1:22 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	6
events	0	constraints	0
reactions	12	function definitions	1
global parameters	26	unit definitions	0
rules	1	initial assignments	0

Model Notes

This a model from the article:

Theoretical and experimental evidence for hysteresis in cell proliferation.

Bai S, Goodrich D, Thron CD, Tecarro E, Obeyesekere M. Cell Cycle. 2003 Jan-Feb;2(1):46-52. [12695688](#) ,

Abstract:

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We propose a mathematical model for the regulation of the G1-phase of the mammalian cell cycle taking into account interactions of cyclin D/cdk4, cyclin E/cdk2, Rb and E2F. Mathematical analysis of this model predicts that a change in the proliferative status in response to a change in concentrations of serum growth factors will exhibit the property of hysteresis: the concentration of growth factors required to induce proliferation is higher than the concentration required to maintain proliferation. We experimentally confirmed this prediction in mouse embryonic fibroblasts in vitro. In agreement with the mathematical model, this indicates that changes in proliferative mode caused by small changes in concentrations of growth factors are not easily reversible. Based on this study, we discuss the importance of proliferation hysteresis for cell cycle regulation.

The original model was taken from the Cell Cycle DataBase (CCDB).

Variable added: assignment rule for denoting phosphorylated Rb (Rb_phosphorylated i.e(RT-RS-R)) created.

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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 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 m^2

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	Size	Unit	Constant	Outside
			Dimensions				
cell	cell		3	1	litre	<input checked="" type="checkbox"/>	

3.1 Compartment `cell`

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

Name `cell`

4 Species

This model contains six species. 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
D_1	D	cell	$\text{mol} \cdot \text{l}^{-1}$	\square	\square
E_1	E	cell	$\text{mol} \cdot \text{l}^{-1}$	\square	\square
R_1	R	cell	$\text{mol} \cdot \text{l}^{-1}$	\square	\square
RS_1	RS	cell	$\text{mol} \cdot \text{l}^{-1}$	\square	\square
theta_1	theta	cell	$\text{mol} \cdot \text{l}^{-1}$	\square	\square
X_1	X	cell	$\text{mol} \cdot \text{l}^{-1}$	\square	\square

5 Parameters

This model contains 26 global parameters.

Table 4: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
GF_1	GF		6.300		<input checked="" type="checkbox"/>
k1_1	k1		0.050		<input checked="" type="checkbox"/>
dD_1	dD		0.400		<input checked="" type="checkbox"/>
aD_1	aD		0.400		<input checked="" type="checkbox"/>
aE_1	aE		0.160		<input checked="" type="checkbox"/>
k2_1	k2		1000.000		<input checked="" type="checkbox"/>
aF_1	aF		0.900		<input checked="" type="checkbox"/>
pX_1	pX		0.480		<input checked="" type="checkbox"/>
RT_1	RT		2.500		<input checked="" type="checkbox"/>
qX_1	qX		0.800		<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"/>
atheta_1	atheta		0.050		<input checked="" type="checkbox"/>
k3_1	k3		1.500		<input checked="" type="checkbox"/>
dtheta_1	dtheta		0.120		<input checked="" type="checkbox"/>
qtheta_1	qtheta		0.300		<input checked="" type="checkbox"/>
aX_1	aX		0.080		<input checked="" type="checkbox"/>
f_1	f		0.350		<input checked="" type="checkbox"/>
g_1	g		0.528		<input checked="" type="checkbox"/>
dX_1	dX		1.040		<input checked="" type="checkbox"/>
dE_1	dE		0.200		<input checked="" type="checkbox"/>
fC_1_1	fe		0.003		<input checked="" type="checkbox"/>
Rb_phos	Rb_phosphorylated		0.000		<input type="checkbox"/>

6 Function definition

This is an overview of one function definition.

6.1 Function definition `Mass_Action_2_1`

Name `Mass_Action_2`

Arguments `k1, S1, S2`

Mathematical Expression

$$k1 \cdot S1 \cdot S2 \quad (1)$$

7 Rule

This is an overview of one rule.

7.1 Rule Rb_phos

Rule Rb_phos is an assignment rule for parameter Rb_phos:

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

8 Reactions

This model contains twelve 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	cyclinD synthesis	$\emptyset \longrightarrow D_1$	
2	cyclinD_1	cyclinD degradation	$D_1 \xrightarrow{D_1, E_1} \emptyset$	
3	cyclinEsynthesis_1	cyclinE synthesis	$\emptyset \xrightarrow{\text{theta_1}} E_1$	
4	cyclinEdegradation_1	cyclinE degradation	$E_1 \xrightarrow{X_1, E_1} \emptyset$	
5	pRBsynthesis_1	pRB synthesis	$\emptyset \xrightarrow{RS_1, R_1, X_1} R_1$	
6	pRBdeplation_1	pRB/E2F complex association	$R_1 + \text{theta_1} \longrightarrow RS_1$	
7	_2	pRB/E2F complex dissociation via cyclin D	$RS_1 \xrightarrow{RS_1, D_1} \text{theta_1}$	
8	null2_1	pRB/E2F complex dissociation via cyclin E	$RS_1 \xrightarrow{RS_1, E_1} \text{theta_1}$	
9	null3_1	E2F synthesis	$\emptyset \xrightarrow{\text{theta_1}} \text{theta_1}$	
10	E2FdegradationviaE2Fdegradation_1	E2F degradation	$\text{theta_1} \xrightarrow{\text{theta_1}, X_1} \emptyset$	
11	cellcycleprogression_1	cell cycle progression go	$\emptyset \xrightarrow{E_1, \text{theta_1}, X_1} X_1$	
12	cellcycleprogression_1	cell cycle progression slow	$X_1 \xrightarrow{X_1} \emptyset$	

8.1 Reaction _1

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

Name cyclinD synthesis

Reaction equation



Product

Table 6: Properties of each product.

Id	Name	SBO
D_1	D	

Kinetic Law

Derived unit contains undeclared units

$$v_1 = \text{vol}(\text{cell}) \cdot aD_1 \cdot \frac{GF_1}{k1_1^1 + GF_1} \quad (4)$$

8.2 Reaction cyclinD_1

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

Name cyclinD 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

$$v_2 = \text{vol}(\text{cell}) \cdot dD_1 \cdot [E_1] \cdot [D_1] \quad (6)$$

8.3 Reaction `cyclinEsynthesis_1`

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

Name cyclinE synthesis

Reaction equation



Modifier

Table 9: Properties of each modifier.

Id	Name	SBO
theta_1	theta	

Product

Table 10: Properties of each product.

Id	Name	SBO
E_1	E	

Kinetic Law

Derived unit contains undeclared units

$$v_3 = \text{vol}(\text{cell}) \cdot aE_1 \cdot \left(\frac{GF_1}{k2_1^I + GF_1} + aF_1 \cdot [\text{theta_1}] \right) \quad (8)$$

8.4 Reaction cyclinEdegradation_1

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

Name cyclinE 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
X_1	X	
E_1	E	

Kinetic Law

Derived unit contains undeclared units

$$v_4 = \text{vol}(\text{cell}) \cdot dE_1 \cdot [X_1] \cdot [E_1] \quad (10)$$

8.5 Reaction pRBsynthesis_1

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

Name pRB synthesis

Reaction equation



Modifiers

Table 13: Properties of each modifier.

Id	Name	SBO
RS_1	RS	
R_1	R	
X_1	X	

Product

Table 14: Properties of each product.

Id	Name	SBO
R_1	R	

Kinetic Law

Derived unit contains undeclared units

$$v_5 = \text{vol}(\text{cell}) \cdot \frac{pX_1 \cdot (RT_1 - [RS_1] - [R_1]) \cdot [X_1]}{qX_1 + RT_1 - [RS_1] - [R_1] + [X_1]} \quad (12)$$

8.6 Reaction `pRBdeplation_1`

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

Name pRB/E2F complex association

Reaction equation



Reactants

Table 15: Properties of each reactant.

Id	Name	SBO
R_1	R	
theta_1	theta	

Product

Table 16: Properties of each product.

Id	Name	SBO
RS_1	RS	

Kinetic Law

Derived unit contains undeclared units

$$v_6 = \text{vol}(\text{cell}) \cdot \text{Mass_Action_2_1}(\text{pS_1}, [\text{R_1}], [\text{theta_1}]) \quad (14)$$

$$\text{Mass_Action_2_1}(k_1, S_1, S_2) = k_1 \cdot S_1 \cdot S_2 \quad (15)$$

$$\text{Mass_Action_2_1}(k_1, S_1, S_2) = k_1 \cdot S_1 \cdot S_2 \quad (16)$$

8.7 Reaction _2

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

Name pRB/E2F complex dissociation via cyclin D

Reaction equation



Reactant

Table 17: Properties of each reactant.

Id	Name	SBO
RS_1	RS	

Modifiers

Table 18: Properties of each modifier.

Id	Name	SBO
RS_1	RS	
D_1	D	

Product

Table 19: Properties of each product.

Id	Name	SBO
theta_1	theta	

Kinetic Law

Derived unit contains undeclared units

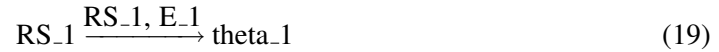
$$v_7 = \text{vol}(\text{cell}) \cdot \frac{pD_1 \cdot [RS_1] \cdot [D_1]}{qD_1 + [RS_1] + [D_1]} \quad (18)$$

8.8 Reaction null2_1

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

Name pRB/E2F complex dissociation via cyclin E

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	

Product

Table 22: Properties of each product.

Id	Name	SBO
theta_1	theta	

Kinetic Law**Derived unit** contains undeclared units

$$v_8 = \text{vol}(\text{cell}) \cdot \frac{pE_1 \cdot [RS_1] \cdot [E_1]}{qE_1 + [RS_1] + [E_1]} \quad (20)$$

8.9 Reaction null3_1

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

Name E2F synthesis**Reaction equation****Modifier**

Table 23: Properties of each modifier.

Id	Name	SBO
theta_1	theta	

Product

Table 24: Properties of each product.

Id	Name	SBO
theta_1	theta	

Kinetic Law**Derived unit** contains undeclared units

$$v_9 = \text{vol}(\text{cell}) \cdot \text{atheta_1} \cdot \left(\frac{GF_1}{k3_1^1 + GF_1} + fC_1_1 \cdot [\text{theta_1}] \right) \quad (22)$$

8.10 Reaction [E2Fdegradationviacellcycleprogression_1](#)

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

Name E2F degradation via cell cycle progression

Reaction equation



Reactant

Table 25: Properties of each reactant.

Id	Name	SBO
theta_1	theta	

Modifiers

Table 26: Properties of each modifier.

Id	Name	SBO
theta_1	theta	
X_1	X	

Kinetic Law

Derived unit contains undeclared units

$$v_{10} = \text{vol}(\text{cell}) \cdot d\text{theta_1} \cdot \frac{[\text{X_1}]}{q\text{theta_1} + [\text{X_1}]} \cdot [\text{theta_1}] \quad (24)$$

8.11 Reaction [cellcycleprogression_1](#)

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

Name cell cycle progression go

Reaction equation



Modifiers

Table 27: Properties of each modifier.

Id	Name	SBO
E_1	E	
theta_1	theta	
X_1	X	

Product

Table 28: Properties of each product.

Id	Name	SBO
X_1	X	

Kinetic Law

Derived unit contains undeclared units

$$v_{11} = \text{vol}(\text{cell}) \cdot (a_{X_1} \cdot [E_1] + f_1 \cdot [\text{theta}_1] + g_1 \cdot [X_1]^2 \cdot [E_1]) \quad (26)$$

8.12 Reaction `cellcycleprogressionslow_1`

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

Name cell cycle progression slow

Reaction equation



Reactant

Table 29: Properties of each reactant.

Id	Name	SBO
X_1	X	

Modifier

Table 30: Properties of each modifier.

Id	Name	SBO
X_1	X	

Kinetic Law

Derived unit contains undeclared units

$$v_{12} = \text{vol}(\text{cell}) \cdot dX_1 \cdot [X_1] \quad (28)$$

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.

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.

9.1 Species D_1

Name D

Initial amount 0 mol

This species takes part in four reactions (as a reactant in [cyclinD_1](#) and as a product in [_1](#) and as a modifier in [cyclinD_1](#), [_2](#)).

$$\frac{d}{dt}D_1 = v_1 - v_2 \quad (29)$$

9.2 Species E_1

Name E

Initial amount 0 mol

This species takes part in six reactions (as a reactant in [cyclinEdegradation_1](#) and as a product in [cyclinEsynthesis_1](#) and as a modifier in [cyclinD_1](#), [cyclinEdegradation_1](#), [null2_1](#), [cellcycleprogression_1](#)).

$$\frac{d}{dt}E_1 = v_3 - v_4 \quad (30)$$

9.3 Species `R_1`

Name `R`

Initial amount 2.5 mol

This species takes part in three reactions (as a reactant in `pRBdeplation_1` and as a product in `pRBsynthesis_1` and as a modifier in `pRBsynthesis_1`).

$$\frac{d}{dt}R_1 = v_5 - v_6 \quad (31)$$

9.4 Species `RS_1`

Name `RS`

Initial amount 0 mol

This species takes part in six reactions (as a reactant in `_2`, `null2_1` and as a product in `pRBdeplation_1` and as a modifier in `pRBsynthesis_1`, `_2`, `null2_1`).

$$\frac{d}{dt}RS_1 = v_6 - v_7 - v_8 \quad (32)$$

9.5 Species `theta_1`

Name `theta`

Initial amount 0 mol

This species takes part in nine reactions (as a reactant in `pRBdeplation_1`, `E2Fdegradationviacellcycleprogression_1` and as a product in `_2`, `null2_1`, `null3_1` and as a modifier in `cyclinEsynthesis_1`, `null3_1`, `E2Fdegradationviacellcycleprogression_1`, `cellcycleprogression_1`).

$$\frac{d}{dt}\theta_1 = v_7 + v_8 + v_9 - v_6 - v_{10} \quad (33)$$

9.6 Species `X_1`

Name `X`

Initial amount 0 mol

This species takes part in seven reactions (as a reactant in `cellcycleprogressionslow_1` and as a product in `cellcycleprogression_1` and as a modifier in `cyclinEdegradation_1`, `pRBsynthesis_1`, `E2Fdegradationviacellcycleprogression_1`, `cellcycleprogression_1`, `cellcycleprogressionslow_1`).

$$\frac{d}{dt}X_1 = v_{11} - v_{12} \quad (34)$$

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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