

SBML Model Report

Model name: “Queralt2006_MitoticExit- _Cdc55DownregulationBySeparase”



May 6, 2016

1 General Overview

This is a document in SBML Level 2 Version 4 format. This model was created by the following two authors: Lukas Endler¹ and Vijayalakshmi Chelliah² at August 20th 2009 at 4:15 p. m. and last time modified at July 16th 2012 at 12:01 a. 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	24
events	0	constraints	0
reactions	31	function definitions	0
global parameters	54	unit definitions	0
rules	10	initial assignments	0

Model Notes

This model is from the article:

Downregulation of PP2A(Cdc55) phosphatase by separase initiates mitotic exit in budding yeast.

Queralt E, Lehane C, Novak B, Uhlmann F. *Cell*. 2006 May 19;125(4):719-32. [16713564](#),

¹EMBL-EBI, lukas@ebi.ac.uk

²EMBL-EBI, viji@ebi.ac.uk

Abstract:

After anaphase, the high mitotic cyclin-dependent kinase (Cdk) activity is downregulated to promote exit from mitosis. To this end, in the budding yeast *S. cerevisiae*, the Cdk counteracting phosphatase Cdc14 is activated. In metaphase, Cdc14 is kept inactive in the nucleolus by its inhibitor Net1. During anaphase, Cdk- and Polo-dependent phosphorylation of Net1 is thought to release active Cdc14. How Net1 is phosphorylated specifically in anaphase, when mitotic kinase activity starts to decline, has remained unexplained. Here, we show that PP2A(Cdc55) phosphatase keeps Net1 underphosphorylated in metaphase. The sister chromatid-separating protease separase, activated at anaphase onset, interacts with and downregulates PP2A(Cdc55), thereby facilitating Cdk-dependent Net1 phosphorylation. PP2A(Cdc55) downregulation also promotes phosphorylation of Bfa1, contributing to activation of the „mitotic exit network,, that sustains Cdc14 as Cdk activity declines. These findings allow us to present a new quantitative model for mitotic exit in budding yeast.

This model originates from BioModels Database: A Database of Annotated Published Models (<http://www.ebi.ac.uk/biomodels/>). It is copyright (c) 2005-2012 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 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
<code>compartment</code>	<code>compartment</code>	0000290	3	1	litre	<input checked="" type="checkbox"/>	

3.1 Compartment `compartment`

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

Name `compartment`

SBO:0000290 physical compartment

4 Species

This model contains 24 species. The boundary condition of eleven 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
AA	AA	compartment	$\text{mol} \cdot \text{l}^{-1}$	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
Clb2	Clb2	compartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
degr	degr	compartment	$\text{mol} \cdot \text{l}^{-1}$	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
Cdc20	Cdc20	compartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
Cdh1	Cdh1	compartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
Cdh1_i	inactive Cdh1	compartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
securinT	securinT	compartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
securin	securin	compartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
separaseT	separaseT	compartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
separase	separase	compartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
securinseparase	securin:separase	compartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
Net1	Net1	compartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
Net1Cdc14	Net1Cdc14	compartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
PoloT	PoloT	compartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
Polo	Polo	compartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
Polo_i	inactive Polo	compartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
Tem1	Tem1	compartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
Tem1_i	inactive Tem1	compartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
Cdc15	Cdc15	compartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
Cdc15_i	inactive Cdc15	compartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
MEN	MEN	compartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>

Id	Name	Compartment	Derived Unit	Constant	Boundary Condi- tion
PP2A	PP2A	compartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
Net1P	Net1P	compartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
Cdc14	Cdc14	compartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>

5 Parameters

This model contains 54 global parameters.

Table 4: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
Cdk	Cdk activity		0.000		<input type="checkbox"/>
ksclb2	ksclb2		0.030		<input checked="" type="checkbox"/>
kdc1b2	kdc1b2		0.030		<input checked="" type="checkbox"/>
kadclb2	kadclb2		0.200		<input checked="" type="checkbox"/>
kaadclb2	kaadclb2		2.000		<input checked="" type="checkbox"/>
kscdc20	kscdc20		0.015		<input checked="" type="checkbox"/>
kdcdc20	kdcdc20		0.050		<input checked="" type="checkbox"/>
kadc20	kadc20		2.000		<input checked="" type="checkbox"/>
kcdh	kcdh		0.010		<input checked="" type="checkbox"/>
kadcdh	kadcdh		1.000		<input checked="" type="checkbox"/>
kapcdh	kapcdh		1.000		<input checked="" type="checkbox"/>
Jcdh	Jcdh		0.002		<input checked="" type="checkbox"/>
Cdh1T	Cdh1T		1.000		<input checked="" type="checkbox"/>
kssecurin	kssecurin		0.030		<input checked="" type="checkbox"/>
kdsecurin	kdsecurin		0.050		<input checked="" type="checkbox"/>
kadsecurin	kadsecurin		2.000		<input checked="" type="checkbox"/>
ksseparase	ksseparase		0.001		<input checked="" type="checkbox"/>
kdseparase	kdseparase		0.004		<input checked="" type="checkbox"/>
lasecurin	lasecurin		500.000		<input checked="" type="checkbox"/>
ldsecurin	ldsecurin		1.000		<input checked="" type="checkbox"/>
kad	kad		0.100		<input checked="" type="checkbox"/>
kd	kd		0.450		<input checked="" type="checkbox"/>
kp	kp		0.400		<input checked="" type="checkbox"/>
kap	kap		2.000		<input checked="" type="checkbox"/>
Jnet	Jnet		0.200		<input checked="" type="checkbox"/>
lanet	lanet		500.000		<input checked="" type="checkbox"/>
ldnet	ldnet		1.000		<input checked="" type="checkbox"/>
Net1T	Net1T		1.000		<input checked="" type="checkbox"/>
Cdc14T	Cdc14T		0.500		<input checked="" type="checkbox"/>
kspolo	kspolo		0.010		<input checked="" type="checkbox"/>
kdpolo	kdpolo		0.010		<input checked="" type="checkbox"/>
kadpolo	kadpolo		0.250		<input checked="" type="checkbox"/>
kapolo	kapolo		0.000		<input checked="" type="checkbox"/>
kaapolo	kaapolo		0.500		<input checked="" type="checkbox"/>
kipolo	kipolo		0.100		<input checked="" type="checkbox"/>
Jpolo	Jpolo		0.250		<input checked="" type="checkbox"/>
katem	katem		0.000		<input checked="" type="checkbox"/>

Id	Name	SBO	Value	Unit	Constant
kaatem	kaatem		0.500		✓
kitem	kitem		0.100		✓
kaitem	kaitem		1.000		✓
Jtem1	Jtem1		0.005		✓
Tem1T	Tem1T		1.000		✓
kacdc15	kacdc15		0.020		✓
kaacdc15	kaacdc15		0.500		✓
kicdc15	kicdc15		0.000		✓
kaicdc15	kaicdc15		0.200		✓
Jcdc15	Jcdc15		0.200		✓
lamen	lamen		10.000		✓
ldmen	ldmen		0.100		✓
Cdc15T	Cdc15T		1.000		✓
kpp	kpp		0.100		✓
ki	ki		20.000		✓
PP2AT	PP2AT		1.000		✓
Inh	Inh		0.000		✓

6 Rules

This is an overview of ten rules.

6.1 Rule `separase`

Rule `separase` is an assignment rule for species `separase`:

$$\text{separase} = [\text{separaseT}] - [\text{securinseparase}] \quad (1)$$

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

6.2 Rule `Cdh1_i`

Rule `Cdh1_i` is an assignment rule for species `Cdh1_i`:

$$\text{Cdh1}_i = \text{Cdh1T} - [\text{Cdh1}] \quad (2)$$

6.3 Rule `Tem1_i`

Rule `Tem1_i` is an assignment rule for species `Tem1_i`:

$$\text{Tem1}_i = \text{Tem1T} - [\text{Tem1}] \quad (3)$$

6.4 Rule securin

Rule securin is an assignment rule for species securin:

$$\text{securin} = [\text{securinT}] - [\text{securinseparase}] \quad (4)$$

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

6.5 Rule Polo_i

Rule Polo_i is an assignment rule for species Polo_i:

$$\text{Polo}_i = [\text{PoloT}] - [\text{Polo}] \quad (5)$$

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

6.6 Rule Cdk

Rule Cdk is an assignment rule for parameter Cdk:

$$\text{Cdk} = \frac{[\text{Clb2}]}{1 + \text{Inh}} \quad (6)$$

6.7 Rule Net1P

Rule Net1P is an assignment rule for species Net1P:

$$\text{Net1P} = \text{Net1T} - [\text{Net1}] - [\text{Net1Cdc14}] \quad (7)$$

6.8 Rule Cdc14

Rule Cdc14 is an assignment rule for species Cdc14:

$$\text{Cdc14} = \text{Cdc14T} - [\text{Net1Cdc14}] \quad (8)$$

6.9 Rule Cdc15_i

Rule Cdc15_i is an assignment rule for species Cdc15_i:

$$\text{Cdc15}_i = \text{Cdc15T} - [\text{Cdc15}] \quad (9)$$

6.10 Rule PP2A

Rule PP2A is an assignment rule for species PP2A:

$$\text{PP2A} = \frac{1 + k_{pp} \cdot k_i \cdot [\text{separase}]}{1 + k_i \cdot [\text{separase}]} \cdot \text{PP2AT} \quad (10)$$

7 Reactions

This model contains 31 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	v1	v1	$AA \rightleftharpoons Clb2$	0000393
2	v2	v2	$Clb2 \xrightarrow{Cdc20, Cdh1} \text{degr}$	0000179
3	v3	v3	$AA \rightleftharpoons Cdc20$	0000393
4	v4	v4	$Cdc20 \xrightarrow{Cdh1} \text{degr}$	0000179
5	v5	v5	$Cdh1_i \xrightarrow{Cdc14} Cdh1$	0000181
6	v6	v6	$Cdh1 \rightleftharpoons Cdh1_i$	0000181
7	v7	v7	$AA \rightleftharpoons \text{securinT} + \text{securin}$	0000393
8	v8	v8	$\text{securinT} + \text{securin} \xrightarrow{Cdc20} \text{degr}$	0000179
9	v9	v9	$AA \rightleftharpoons \text{separaseT} + \text{separase}$	0000393
10	v10	v10	$\text{separaseT} + \text{separase} \rightleftharpoons \text{degr}$	0000179
11	v11	v11	$\text{securin} + \text{separase} \rightleftharpoons \text{securinseparase}$	0000526
12	securinseparase- _securindegradation	securinseparase securindegradation	$\text{securinseparase} \xrightarrow{Cdc20} \text{separase}$	0000179
13	securinseparase- _separase- _degradation	securin:separase separase degradation	$\text{securinseparase} \rightleftharpoons \text{securin}$	0000179
14	v13	Net1P dephosphorylation	$\text{Net1P} \xrightarrow{Cdc14, Clb2, PP2A} \text{Net1}$	0000330
15	v14	Net1 phosphorylation	$\text{Net1} \xrightarrow{MEN, Net1Cdc14, Clb2} \text{Net1P}$	0000216
16	v15	v15	$\text{Net1} \xrightarrow{Cdc14} \text{Net1Cdc14}$	0000526

Nº	Id	Name	Reaction Equation	SBO
17	v16	v16	$\text{Net1Cdc14} \rightleftharpoons \text{Net1}$	0000180
18	v17	Net1Cdc14 phosphorylation	$\text{Net1Cdc14} \xrightleftharpoons{\text{MEN, Net1, Clb2}} \text{Net1P}$	0000216
19	v18	v18	$\text{AA} \rightleftharpoons \text{PoloT} + \text{Polo.i}$	0000393
20	v19	v19	$\text{PoloT} + \text{Polo.i} \xrightleftharpoons{\text{Cdh1}} \text{degr}$	0000179
21	v20	v20	$\text{Polo.i} \xrightleftharpoons{\text{PoloT}} \text{Polo}$	0000181
22	v21	v21	$\text{Polo} \rightleftharpoons \text{Polo.i}$	0000181
23	v22	v22	$\text{Polo} \xrightleftharpoons{\text{Cdh1}} \text{degr}$	0000179
24	v23	v23	$\text{Tem1.i} \xrightleftharpoons{\text{Polo}} \text{Tem1}$	0000181
25	v24	v24	$\text{Tem1} \xrightleftharpoons{\text{PP2A}} \text{Tem1.i}$	0000181
26	v25	v25	$\text{Cdc15.i} \xrightleftharpoons{\text{Cdc14}} \text{Cdc15}$	0000181
27	v26	v26	$\text{Cdc15} \rightleftharpoons \text{Cdc15.i}$	0000181
28	v27	v27	$\text{AA} \xrightleftharpoons{\text{Tem1, Cdc15}} \text{MEN}$	0000393
29	v28	v28	$\text{MEN} \rightleftharpoons \text{degr}$	0000179
30	v29	Tem1 degradation in MEN	$\text{MEN} \xrightleftharpoons{\text{Tem1, PP2A}} \emptyset$	0000179
31	v30	Cdc15 degradation in MEN	$\text{MEN} \xrightleftharpoons{\text{Cdc15, Clb2}} \emptyset$	0000179

7.1 Reaction v1

This is a reversible reaction of one reactant forming one product.

Name v1

SBO:0000393 production

Reaction equation



Reactant

Table 6: Properties of each reactant.

Id	Name	SBO
AA	AA	

Product

Table 7: Properties of each product.

Id	Name	SBO
Clb2	Clb2	

Kinetic Law

Derived unit not available

$v_1 = k_{\text{sclb2}}$

(12)

7.2 Reaction v2

This is a reversible reaction of one reactant forming one product influenced by two modifiers.

Name v2

SBO:0000179 degradation

Reaction equation



Reactant

Table 8: Properties of each reactant.

Id	Name	SBO
C1b2	Clb2	

Modifiers

Table 9: Properties of each modifier.

Id	Name	SBO
Cdc20	Cdc20	
Cdh1	Cdh1	

Product

Table 10: Properties of each product.

Id	Name	SBO
degr	degr	

Kinetic Law

Derived unit contains undeclared units

$$v_2 = (kdclb2 + kadclb2 \cdot [Cdc20] + kaadclb2 \cdot [Cdh1]) \cdot [Clb2] \quad (14)$$

7.3 Reaction v3

This is a reversible reaction of one reactant forming one product.

Name v3

SBO:0000393 production

Reaction equation



Reactant

Table 11: Properties of each reactant.

Id	Name	SBO
AA	AA	

Product

Table 12: Properties of each product.

Id	Name	SBO
Cdc20	Cdc20	

Kinetic Law

Derived unit not available

$$v_3 = kscdc20 \quad (16)$$

7.4 Reaction v4

This is a reversible reaction of one reactant forming one product influenced by one modifier.

Name v4

SBO:0000179 degradation

Reaction equation



Reactant

Table 13: Properties of each reactant.

Id	Name	SBO
Cdc20	Cdc20	

Modifier

Table 14: Properties of each modifier.

Id	Name	SBO
Cdh1	Cdh1	

Product

Table 15: Properties of each product.

Id	Name	SBO
degr	degr	

Kinetic Law

Derived unit contains undeclared units

$$v_4 = (k_{dc20} + k_{ad20} \cdot [Cdh1]) \cdot [Cdc20] \quad (18)$$

7.5 Reaction v5

This is a reversible reaction of one reactant forming one product influenced by one modifier.

Name v5

SBO:0000181 conformational transition

Reaction equation



Reactant

Table 16: Properties of each reactant.

Id	Name	SBO
Cdh1_i	inactive Cdh1	

Modifier

Table 17: Properties of each modifier.

Id	Name	SBO
Cdc14	Cdc14	

Product

Table 18: Properties of each product.

Id	Name	SBO
Cdh1	Cdh1	

Kinetic Law

Derived unit contains undeclared units

$$v_5 = (k_{dcdh} + k_{adcdh} \cdot [Cdc14]) \cdot \frac{Cdh1T - [Cdh1]}{J_{cdh} + Cdh1T - [Cdh1]} \quad (20)$$

7.6 Reaction v6

This is a reversible reaction of one reactant forming one product.

Name v6

SBO:0000181 conformational transition

Reaction equation



Reactant

Table 19: Properties of each reactant.

Id	Name	SBO
Cdh1	Cdh1	

Product

Table 20: Properties of each product.

Id	Name	SBO
Cdh1_i	inactive Cdh1	

Kinetic Law

Derived unit contains undeclared units

$$v_6 = \text{kapcdh} \cdot \text{Cdk} \cdot \frac{[\text{Cdh1}]}{\text{Jcdh} + [\text{Cdh1}]}$$

(22)

7.7 Reaction v7

This is a reversible reaction of one reactant forming two products.

Name v7

SBO:0000393 production

Reaction equation



Reactant

Table 21: Properties of each reactant.

Id	Name	SBO
AA	AA	

Products

Table 22: Properties of each product.

Id	Name	SBO
securinT	securinT	
securin	securin	

Kinetic Law

Derived unit not available

$$v_7 = k_{\text{securin}} \quad (24)$$

7.8 Reaction v8

This is a reversible reaction of two reactants forming one product influenced by one modifier.

Name v8

SBO:0000179 degradation

Reaction equation



Reactants

Table 23: Properties of each reactant.

Id	Name	SBO
securinT	securinT	
securin	securin	

Modifier

Table 24: Properties of each modifier.

Id	Name	SBO
Cdc20	Cdc20	

Product

Table 25: Properties of each product.

Id	Name	SBO
degr	degr	

Kinetic Law

Derived unit contains undeclared units

$$v_8 = (k_{\text{dsecurin}} + k_{\text{adsecurin}} \cdot [\text{Cdc20}]) \cdot [\text{securinT}] \quad (26)$$

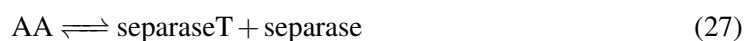
7.9 Reaction v9

This is a reversible reaction of one reactant forming two products.

Name v9

SBO:0000393 production

Reaction equation



Reactant

Table 26: Properties of each reactant.

Id	Name	SBO
AA	AA	

Products

Table 27: Properties of each product.

Id	Name	SBO
separaseT	separaseT	
separase	separase	

Kinetic Law

Derived unit not available

$$v_9 = k_{\text{separase}} \quad (28)$$

7.10 Reaction v10

This is a reversible reaction of two reactants forming one product.

Name v10

SBO:0000179 degradation

Reaction equation



Reactants

Table 28: Properties of each reactant.

Id	Name	SBO
separaseT	separaseT	
separase	separase	

Product

Table 29: Properties of each product.

Id	Name	SBO
degr	degr	

Kinetic Law

Derived unit contains undeclared units

$$v_{10} = kdseparase \cdot [separaseT] \quad (30)$$

7.11 Reaction v11

This is a reversible reaction of two reactants forming one product.

Name v11

SBO:0000526 protein complex formation

Reaction equation



Reactants

Table 30: Properties of each reactant.

Id	Name	SBO
securin	securin	
separase	separase	

Product

Table 31: Properties of each product.

Id	Name	SBO
securinseparase	securin:separase	

Kinetic Law

Derived unit contains undeclared units

$$v_{11} = lasecurin \cdot [securin] \cdot [separase] - ldsecurin \cdot [securinseparase] \quad (32)$$

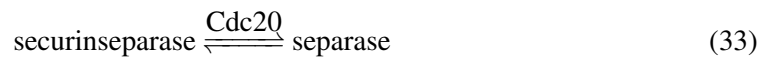
7.12 Reaction securinseparase.securindegradation

This is a reversible reaction of one reactant forming one product influenced by one modifier.

Name securinseparase securindegradation

SBO:0000179 degradation

Reaction equation



Reactant

Table 32: Properties of each reactant.

Id	Name	SBO
securinseparase	securin:separase	

Modifier

Table 33: Properties of each modifier.

Id	Name	SBO
Cdc20	Cdc20	

Product

Table 34: Properties of each product.

Id	Name	SBO
separase	separase	

Kinetic Law

Derived unit contains undeclared units

$$v_{12} = (kd_{\text{securin}} + ka_{\text{securin}} \cdot [\text{Cdc20}]) \cdot [\text{securinseparase}] \quad (34)$$

7.13 Reaction securinseparase.separase.degradation

This is a reversible reaction of one reactant forming one product.

Name securin:separase separase degradation

SBO:0000179 degradation

Reaction equation



Reactant

Table 35: Properties of each reactant.

Id	Name	SBO
securinseparase	securin:separase	

Product

Table 36: Properties of each product.

Id	Name	SBO
securin	securin	

Kinetic Law

Derived unit contains undeclared units

$$v_{13} = kd_{\text{separase}} \cdot [\text{securinseparase}] \quad (36)$$

7.14 Reaction v13

This is a reversible reaction of one reactant forming one product influenced by three modifiers.

Name Net1P dephosphorylation

SBO:0000330 dephosphorylation

Reaction equation



Reactant

Table 37: Properties of each reactant.

Id	Name	SBO
Net1P	Net1P	

Modifiers

Table 38: Properties of each modifier.

Id	Name	SBO
Cdc14	Cdc14	
Clb2	Clb2	
PP2A	PP2A	

Product

Table 39: Properties of each product.

Id	Name	SBO
Net1	Net1	

Kinetic Law

Derived unit contains undeclared units

$$v_{14} = (\text{kad} \cdot [\text{Cdc14}] + \text{kd} \cdot [\text{PP2A}]) \cdot \frac{[\text{Net1P}]}{\text{Jnet} + [\text{Net1P}]} \quad (38)$$

7.15 Reaction v14

This is a reversible reaction of one reactant forming one product influenced by three modifiers.

Name Net1 phosphorylation

SBO:0000216 phosphorylation

Reaction equation



Reactant

Table 40: Properties of each reactant.

Id	Name	SBO
Net1	Net1	

Modifiers

Table 41: Properties of each modifier.

Id	Name	SBO
MEN	MEN	
Net1Cdc14	Net1Cdc14	
Clb2	Clb2	

Product

Table 42: Properties of each product.

Id	Name	SBO
Net1P	Net1P	

Kinetic Law

Derived unit contains undeclared units

$$v_{15} = (\text{kp} \cdot \text{Cdk} + \text{kap} \cdot [\text{MEN}]) \cdot \frac{[\text{Net1}]}{\text{jnet} + [\text{Net1}] + [\text{Net1Cdc14}]} \quad (40)$$

7.16 Reaction v15

This is a reversible reaction of one reactant forming one product influenced by one modifier.

Name v15

SBO:0000526 protein complex formation

Reaction equation



Reactant

Table 43: Properties of each reactant.

Id	Name	SBO
Net1	Net1	

Modifier

Table 44: Properties of each modifier.

Id	Name	SBO
Cdc14	Cdc14	

Product

Table 45: Properties of each product.

Id	Name	SBO
Net1Cdc14	Net1Cdc14	

Kinetic Law

Derived unit contains undeclared units

$$v_{16} = \text{lanet} \cdot [\text{Net1}] \cdot [\text{Cdc14}] \quad (42)$$

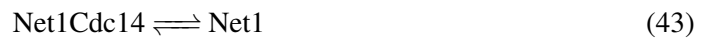
7.17 Reaction v16

This is a reversible reaction of one reactant forming one product.

Name v16

SBO:0000180 dissociation

Reaction equation



Reactant

Table 46: Properties of each reactant.

Id	Name	SBO
Net1Cdc14	Net1Cdc14	

Product

Table 47: Properties of each product.

Id	Name	SBO
Net1	Net1	

Kinetic Law

Derived unit contains undeclared units

$$v_{17} = \text{ldnet} \cdot [\text{Net1Cdc14}] \quad (44)$$

7.18 Reaction v17

This is a reversible reaction of one reactant forming one product influenced by three modifiers.

Name Net1Cdc14 phosphorylation

SBO:0000216 phosphorylation

Reaction equation



Reactant

Table 48: Properties of each reactant.

Id	Name	SBO
Net1Cdc14	Net1Cdc14	

Modifiers

Table 49: Properties of each modifier.

Id	Name	SBO
MEN	MEN	
Net1	Net1	
Clb2	Clb2	

Product

Table 50: Properties of each product.

Id	Name	SBO
Net1P	Net1P	

Kinetic Law

Derived unit contains undeclared units

$$v_{18} = (k_p \cdot Cdk + k_{ap} \cdot [MEN]) \cdot \frac{[Net1Cdc14]}{J_{net} + [Net1] + [Net1Cdc14]} \tag{46}$$

7.19 Reaction v18

This is a reversible reaction of one reactant forming two products.

Name v18

SBO:0000393 production

Reaction equation



Reactant

Table 51: Properties of each reactant.

Id	Name	SBO
AA	AA	

Products

Table 52: Properties of each product.

Id	Name	SBO
PoloT	PoloT	
Polo_i	inactive Polo	

Kinetic Law

Derived unit not available

$$v_{19} = k_{\text{spolo}} \quad (48)$$

7.20 Reaction v19

This is a reversible reaction of two reactants forming one product influenced by one modifier.

Name v19

SBO:0000179 degradation

Reaction equation



Reactants

Table 53: Properties of each reactant.

Id	Name	SBO
PoloT	PoloT	
Polo_i	inactive Polo	

Modifier

Table 54: Properties of each modifier.

Id	Name	SBO
Cdh1	Cdh1	

Product

Table 55: Properties of each product.

Id	Name	SBO
degr	degr	

Kinetic Law

Derived unit contains undeclared units

$$v_{20} = (kdpolo + kadpolo \cdot [Cdh1]) \cdot [PoloT] \quad (50)$$

7.21 Reaction v_{20}

This is a reversible reaction of one reactant forming one product influenced by one modifier.

Name v_{20}

SBO:0000181 conformational transition

Reaction equation



Reactant

Table 56: Properties of each reactant.

Id	Name	SBO
Polo_i	inactive Polo	

Modifier

Table 57: Properties of each modifier.

Id	Name	SBO
Po1oT	PoloT	

Product

Table 58: Properties of each product.

Id	Name	SBO
Po1o	Polo	

Kinetic Law

Derived unit contains undeclared units

$$v_{21} = (\text{kapolo} + \text{kaapolo} \cdot \text{Cdk}) \cdot \frac{[\text{PoloT}] - [\text{Polo}]}{J_{\text{polo}} + [\text{PoloT}] - [\text{Polo}]} \quad (52)$$

7.22 Reaction v21

This is a reversible reaction of one reactant forming one product.

Name v21

SBO:0000181 conformational transition

Reaction equation



Reactant

Table 59: Properties of each reactant.

Id	Name	SBO
Po1o	Polo	

Product

Table 60: Properties of each product.

Id	Name	SBO
Polo_i	inactive Polo	

Kinetic Law

Derived unit contains undeclared units

$$v_{22} = k_{\text{ipolo}} \cdot \frac{[\text{Polo}]}{J_{\text{polo}} + [\text{Polo}]} \tag{54}$$

7.23 Reaction v22

This is a reversible reaction of one reactant forming one product influenced by one modifier.

Name v22

SBO:0000179 degradation

Reaction equation



Reactant

Table 61: Properties of each reactant.

Id	Name	SBO
Polo	Polo	

Modifier

Table 62: Properties of each modifier.

Id	Name	SBO
Cdh1	Cdh1	

Product

Table 63: Properties of each product.

Id	Name	SBO
degr	degr	

Kinetic Law

Derived unit contains undeclared units

$$v_{23} = (kdpolo + kadpolo \cdot [Cdh1]) \cdot [Polo] \quad (56)$$

7.24 Reaction v23

This is a reversible reaction of one reactant forming one product influenced by one modifier.

Name v23

SBO:0000181 conformational transition

Reaction equation



Reactant

Table 64: Properties of each reactant.

Id	Name	SBO
Tem1_i	inactive Tem1	

Modifier

Table 65: Properties of each modifier.

Id	Name	SBO
Polo	Polo	

Product

Table 66: Properties of each product.

Id	Name	SBO
Tem1	Tem1	

Kinetic Law

Derived unit contains undeclared units

$$v_{24} = (k_{\text{atem}} + k_{\text{aatem}} \cdot [\text{Polo}]) \cdot \frac{\text{Tem1T} - [\text{Tem1}]}{J_{\text{tem1}} + \text{Tem1T} - [\text{Tem1}]}$$

(58)

7.25 Reaction v24

This is a reversible reaction of one reactant forming one product influenced by one modifier.

Name v24

SBO:0000181 conformational transition

Reaction equation



Reactant

Table 67: Properties of each reactant.

Id	Name	SBO
Tem1	Tem1	

Modifier

Table 68: Properties of each modifier.

Id	Name	SBO
PP2A	PP2A	

Product

Table 69: Properties of each product.

Id	Name	SBO
Tem1_i	inactive Tem1	

Kinetic Law

Derived unit contains undeclared units

$$v_{25} = (k_{\text{item}} + k_{\text{aitem}} \cdot [\text{PP2A}]) \cdot \frac{[\text{Tem1}]}{J_{\text{tem1}} + [\text{Tem1}]} \quad (60)$$

7.26 Reaction v25

This is a reversible reaction of one reactant forming one product influenced by one modifier.

Name v25

SBO:0000181 conformational transition

Reaction equation



Reactant

Table 70: Properties of each reactant.

Id	Name	SBO
Cdc15_i	inactive Cdc15	

Modifier

Table 71: Properties of each modifier.

Id	Name	SBO
Cdc14	Cdc14	

Product

Table 72: Properties of each product.

Id	Name	SBO
Cdc15	Cdc15	

Kinetic Law

Derived unit contains undeclared units

$$v_{26} = (kacdc15 + kaacdc15 \cdot [Cdc14]) \cdot \frac{Cdc15T - [Cdc15]}{Jcdc15 + Cdc15T - [Cdc15]} \tag{62}$$

7.27 Reaction v26

This is a reversible reaction of one reactant forming one product.

Name v26

SBO:0000181 conformational transition

Reaction equation



Reactant

Table 73: Properties of each reactant.

Id	Name	SBO
Cdc15	Cdc15	

Product

Table 74: Properties of each product.

Id	Name	SBO
Cdc15_i	inactive Cdc15	

Kinetic Law

Derived unit contains undeclared units

$$v_{27} = (kicdc15 + kaicdc15 \cdot Cdk) \cdot \frac{[Cdc15]}{Jcdc15 + [Cdc15]} \tag{64}$$

7.28 Reaction v27

This is a reversible reaction of one reactant forming one product influenced by two modifiers.

Name v27

SBO:0000393 production

Reaction equation



Reactant

Table 75: Properties of each reactant.

Id	Name	SBO
AA	AA	

Modifiers

Table 76: Properties of each modifier.

Id	Name	SBO
Tem1	Tem1	
Cdc15	Cdc15	

Product

Table 77: Properties of each product.

Id	Name	SBO
MEN	MEN	

Kinetic Law

Derived unit contains undeclared units

$$v_{28} = \text{lamen} \cdot ([\text{Tem1}] - [\text{MEN}]) \cdot ([\text{Cdc15}] - [\text{MEN}]) \quad (66)$$

7.29 Reaction v28

This is a reversible reaction of one reactant forming one product.

Name v28

SBO:0000179 degradation

Reaction equation



Reactant

Table 78: Properties of each reactant.

Id	Name	SBO
MEN	MEN	

Product

Table 79: Properties of each product.

Id	Name	SBO
degr	degr	

Kinetic Law

Derived unit contains undeclared units

$v_{29} = \text{Idmen} \cdot [\text{MEN}]$

(68)

7.30 Reaction v29

This is a reversible reaction of one reactant forming no product influenced by two modifiers.

Name Tem1 degradation in MEN

SBO:0000179 degradation

Reaction equation



Reactant

Table 80: Properties of each reactant.

Id	Name	SBO
MEN	MEN	

Modifiers

Table 81: Properties of each modifier.

Id	Name	SBO
Tem1	Tem1	
PP2A	PP2A	

Kinetic Law

Derived unit contains undeclared units

$$v_{30} = (k_{\text{item}} + k_{\text{aitem}} \cdot [\text{PP2A}]) \cdot \frac{[\text{MEN}]}{J_{\text{tem1}} + [\text{Tem1}]} \quad (70)$$

7.31 Reaction v30

This is a reversible reaction of one reactant forming no product influenced by two modifiers.

Name Cdc15 degradation in MEN

SBO:0000179 degradation

Reaction equation



Reactant

Table 82: Properties of each reactant.

Id	Name	SBO
MEN	MEN	

Modifiers

Table 83: Properties of each modifier.

Id	Name	SBO
Cdc15	Cdc15	
Clb2	Clb2	

Kinetic Law

Derived unit contains undeclared units

$$v_{31} = (\text{kidc15} + \text{kaicdc15} \cdot \text{Cdk}) \cdot \frac{[\text{MEN}]}{\text{Jcdc15} + [\text{Cdc15}]} \quad (72)$$

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 AA

Name AA

SBO:0000291 empty set

Initial concentration 1 mol · l⁻¹

This species takes part in six reactions (as a reactant in [v1](#), [v3](#), [v7](#), [v9](#), [v18](#), [v27](#)), which do not influence its rate of change because this constant species is on the boundary of the reaction system:

$$\frac{d}{dt} \text{AA} = 0 \quad (73)$$

8.2 Species Clb2

Name Clb2

SBO:0000252 polypeptide chain

Initial concentration $0.997 \text{ mol} \cdot \text{l}^{-1}$

This species takes part in six reactions (as a reactant in [v2](#) and as a product in [v1](#) and as a modifier in [v13](#), [v14](#), [v17](#), [v30](#)).

$$\frac{d}{dt} \text{Clb2} = v_1 - v_2 \quad (74)$$

8.3 Species [degr](#)

Name [degr](#)

SBO:0000291 empty set

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

This species takes part in seven reactions (as a product in [v2](#), [v4](#), [v8](#), [v10](#), [v19](#), [v22](#), [v28](#)), which do not influence its rate of change because this constant species is on the boundary of the reaction system:

$$\frac{d}{dt} \text{degr} = 0 \quad (75)$$

8.4 Species [Cdc20](#)

Name [Cdc20](#)

SBO:0000252 polypeptide chain

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

This species takes part in five reactions (as a reactant in [v4](#) and as a product in [v3](#) and as a modifier in [v2](#), [v8](#), [securinseparase_securindegradation](#)).

$$\frac{d}{dt} \text{Cdc20} = v_3 - v_4 \quad (76)$$

8.5 Species [Cdh1](#)

Name [Cdh1](#)

SBO:0000252 polypeptide chain

Initial concentration $3.8 \cdot 10^{-5} \text{ mol} \cdot \text{l}^{-1}$

This species takes part in six reactions (as a reactant in [v6](#) and as a product in [v5](#) and as a modifier in [v2](#), [v4](#), [v19](#), [v22](#)).

$$\frac{d}{dt} \text{Cdh1} = v_5 - v_6 \quad (77)$$

8.6 Species `Cdh1_i`

Name inactive Cdh1

SBO:0000252 polypeptide chain

Involved in rule `Cdh1_i`

This species takes part in two reactions (as a reactant in `v5` and as a product in `v6`). Not these but one rule determines the species' quantity because this species is on the boundary of the reaction system.

8.7 Species `securinT`

Name securinT

SBO:0000252 polypeptide chain

Initial concentration $0.6 \text{ mol} \cdot \text{l}^{-1}$

This species takes part in two reactions (as a reactant in `v8` and as a product in `v7`).

$$\frac{d}{dt}\text{securinT} = v_7 - v_8 \quad (78)$$

8.8 Species `securin`

Name securin

SBO:0000252 polypeptide chain

Involved in rule `securin`

This species takes part in four reactions (as a reactant in `v8`, `v11` and as a product in `v7`, `securinseparase_separase_degradation`). Not these but one rule determines the species' quantity because this species is on the boundary of the reaction system.

8.9 Species `separaseT`

Name separaseT

SBO:0000252 polypeptide chain

Initial concentration $0.25 \text{ mol} \cdot \text{l}^{-1}$

This species takes part in two reactions (as a reactant in `v10` and as a product in `v9`).

$$\frac{d}{dt}\text{separaseT} = v_9 - v_{10} \quad (79)$$

8.10 Species *separase*

Name *separase*

SBO:0000252 polypeptide chain

Involved in rule *separase*

This species takes part in four reactions (as a reactant in *v10*, *v11* and as a product in *v9*, *securinseparase_securindegradation*). Not these but one rule determines the species' quantity because this species is on the boundary of the reaction system.

8.11 Species *securinseparase*

Name *securin:separase*

SBO:0000297 protein complex

Initial concentration $0.248 \text{ mol} \cdot \text{l}^{-1}$

This species takes part in three reactions (as a reactant in *securinseparase_securindegradation*, *securinseparase_separase_degradation* and as a product in *v11*).

$$\frac{d}{dt} \text{securinseparase} = v_{11} - v_{12} - v_{13} \quad (80)$$

8.12 Species *Net1*

Name *Net1*

SBO:0000252 polypeptide chain

Initial concentration $0.098 \text{ mol} \cdot \text{l}^{-1}$

This species takes part in five reactions (as a reactant in *v14*, *v15* and as a product in *v13*, *v16* and as a modifier in *v17*).

$$\frac{d}{dt} \text{Net1} = v_{14} + v_{17} - v_{15} - v_{16} \quad (81)$$

8.13 Species *Net1Cdc14*

Name *Net1Cdc14*

SBO:0000297 protein complex

Initial concentration $0.485 \text{ mol} \cdot \text{l}^{-1}$

This species takes part in four reactions (as a reactant in *v16*, *v17* and as a product in *v15* and as a modifier in *v14*).

$$\frac{d}{dt} \text{Net1Cdc14} = v_{16} - v_{17} - v_{18} \quad (82)$$

8.14 Species PoloT

Name PoloT

SBO:0000252 polypeptide chain

Initial concentration $0.99 \text{ mol} \cdot \text{l}^{-1}$

This species takes part in three reactions (as a reactant in v19 and as a product in v18 and as a modifier in v20).

$$\frac{d}{dt}\text{PoloT} = v_{19} - v_{20} \quad (83)$$

8.15 Species Polo

Name Polo

SBO:0000252 polypeptide chain

Initial concentration $0.945 \text{ mol} \cdot \text{l}^{-1}$

This species takes part in four reactions (as a reactant in v21, v22 and as a product in v20 and as a modifier in v23).

$$\frac{d}{dt}\text{Polo} = v_{21} - v_{22} - v_{23} \quad (84)$$

8.16 Species Polo_i

Name inactive Polo

SBO:0000252 polypeptide chain

Involved in rule Polo_i

This species takes part in four reactions (as a reactant in v19, v20 and as a product in v18, v21). Not these but one rule determines the species' quantity because this species is on the boundary of the reaction system.

8.17 Species Tem1

Name Tem1

SBO:0000252 polypeptide chain

Initial concentration $0.00389 \text{ mol} \cdot \text{l}^{-1}$

This species takes part in four reactions (as a reactant in v24 and as a product in v23 and as a modifier in v27, v29).

$$\frac{d}{dt}\text{Tem1} = v_{24} - v_{25} \quad (85)$$

8.18 Species [Tem1_i](#)

Name inactive Tem1

SBO:0000252 polypeptide chain

Involved in rule [Tem1_i](#)

This species takes part in two reactions (as a reactant in [v23](#) and as a product in [v24](#)). Not these but one rule determines the species' quantity because this species is on the boundary of the reaction system.

8.19 Species [Cdc15](#)

Name Cdc15

SBO:0000252 polypeptide chain

Initial concentration $0.0257 \text{ mol} \cdot \text{l}^{-1}$

This species takes part in four reactions (as a reactant in [v26](#) and as a product in [v25](#) and as a modifier in [v27](#), [v30](#)).

$$\frac{d}{dt} \text{Cdc15} = v_{26} - v_{27} \quad (86)$$

8.20 Species [Cdc15_i](#)

Name inactive Cdc15

SBO:0000252 polypeptide chain

Involved in rule [Cdc15_i](#)

This species takes part in two reactions (as a reactant in [v25](#) and as a product in [v26](#)). Not these but one rule determines the species' quantity because this species is on the boundary of the reaction system.

8.21 Species [MEN](#)

Name MEN

SBO:0000297 protein complex

Initial concentration $8 \cdot 10^{-6} \text{ mol} \cdot \text{l}^{-1}$

This species takes part in six reactions (as a reactant in [v28](#), [v29](#), [v30](#) and as a product in [v27](#) and as a modifier in [v14](#), [v17](#)).

$$\frac{d}{dt} \text{MEN} = v_{28} - v_{29} - v_{30} - v_{31} \quad (87)$$

8.22 Species [PP2A](#)

Name PP2A

SBO:0000252 polypeptide chain

Involved in rule [PP2A](#)

This species takes part in three reactions (as a modifier in [v13](#), [v24](#), [v29](#)). Not these but one rule determines the species' quantity because this species is on the boundary of the reaction system.

8.23 Species [Net1P](#)

Name Net1P

SBO:0000297 protein complex

Involved in rule [Net1P](#)

This species takes part in three reactions (as a reactant in [v13](#) and as a product in [v14](#), [v17](#)). Not these but one rule determines the species' quantity because this species is on the boundary of the reaction system.

8.24 Species [Cdc14](#)

Name Cdc14

SBO:0000252 polypeptide chain

Involved in rule [Cdc14](#)

This species takes part in four reactions (as a modifier in [v5](#), [v13](#), [v15](#), [v25](#)). 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:0000179 degradation: Complete disappearance of a physical entity

SBO:0000180 dissociation: Transformation of a non-covalent complex that results in the formation of several independent biochemical entities

SBO:0000181 conformational transition: Biochemical reaction that does not result in the modification of covalent bonds of reactants, but rather modifies the conformation of some reactants, that is the relative position of their atoms in space

SBO:0000216 phosphorylation: Addition of a phosphate group ($\text{-H}_2\text{PO}_4$) to a chemical 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:0000290 physical compartment:** Specific location of space, that can be bounded or not. A physical compartment can have 1, 2 or 3 dimensions
- SBO:0000291 empty set:** Entity defined by the absence of any actual object. An empty set is often used to represent the source of a creation process or the result of a degradation process.
- SBO:0000297 protein complex:** Macromolecular complex containing one or more polypeptide chains possibly associated with simple chemicals. CHEBI:3608
- SBO:0000330 dephosphorylation:** Removal of a phosphate group ($\text{-H}_2\text{PO}_4$) from a chemical entity.
- SBO:0000393 production:** Generation of a material or conceptual entity.
- SBO:0000526 protein complex formation:** The process by which two or more proteins interact non-covalently to form a protein complex (SBO:0000297)

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.

^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