

SBML Model Report

Model name: “Suh2004_KCNQ_Regulation”



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 November ninth 2006 at 6:25 p. m. and last time modified at July fifth 2012 at 4:29 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	3
species types	0	species	23
events	2	constraints	0
reactions	15	function definitions	0
global parameters	12	unit definitions	13
rules	9	initial assignments	0

Model Notes

The model reproduces FIG 11A and FIG 11B of the paper. However, please note that FIG 11B is a plot of normalised amounts versus time. The „stoichiometry„ field has been used to convert fluxes from membrane species to volume species. The value of 0.0009967 is a product of (Surface to Volume_M*(1/Avagadro’s number)*1E21. 0.6 is the surface to volume ratio of the plasma membrane, 1E21 is required for a unit surface to volume ratio and the Avagadro’s number is present in the denominator to convert molecules to moles. The model was successfully tested using MathSBML and SBML ODESolver.

¹California Institute of Technology, hdharuri@cds.caltech.edu

All the kinetic laws have the unit items per second , which requires the one reaction taking place in the cytoplasm - IP3Phosphatase - to include an explicit conversion factor both in the kinetic law and the stoichiometry of IP3_C . The kinetic law is multiplied and the stoichiometry divided by the number of molecules per micro-mole. This conversion factor is only required for correct units and can be replaced by 1, if it should lead to numerical problems.

To the extent possible under law, all copyright and related or neighbouring rights to this encoded model have been dedicated to the public domain worldwide. Please refer to [CC0 Public Domain Dedication](#) for more information.

In summary, you are entitled to use this encoded model in absolutely any manner you deem suitable, verbatim, or with modification, alone or embedded it in a larger context, redistribute it, commercially or not, in a restricted way or not.

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 17 unit definitions of which four are predefined by SBML and not mentioned in the model.

2.1 Unit `micromole`

Definition μmol

2.2 Unit `substance`

Name `items`

Definition `item`

2.3 Unit `items_per_micromole`

Definition $\text{item} \cdot \mu\text{mol}^{-1}$

2.4 Unit `molecules`

Definition `item`

2.5 Unit `um2`

Definition μm^2

2.6 Unit `molecules_um_2_s_1`

Definition $\mu\text{m}^{-2} \cdot \text{s}^{-1} \cdot \text{item}$

2.7 Unit `molecules_um_2`

Definition $\mu\text{m}^{-2} \cdot \text{item}$

2.8 Unit `uM_s_1`

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

2.9 Unit `s_1`

Definition s^{-1}

2.10 Unit `um2_molecules_1_s_1`

Definition $\text{item}^{-1} \cdot \mu\text{m}^2 \cdot \text{s}^{-1}$

2.11 Unit `uM_1_s_1`

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

2.12 Unit `uM_um_s_1`

Definition $\mu\text{m} \cdot \mu\text{mol} \cdot \text{s}^{-1} \cdot \text{l}^{-1}$

2.13 Unit `microMolar`

Name `microMolar`

Definition $\mu\text{mol} \cdot \text{l}^{-1}$

2.14 Unit `volume`

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

Definition `l`

2.15 Unit `area`

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

Definition m^2

2.16 Unit length

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

Definition m

2.17 Unit time

Notes Second is the predefined SBML unit for time.

Definition s

3 Compartments

This model contains three compartments.

Table 2: Properties of all compartments.

Id	Name	SBO	Spatial Dimensions	Size	Unit	Constant	Outside
EX	Extracellular		3	1		<input checked="" type="checkbox"/>	
M	Membrane		2	1	μm^2	<input checked="" type="checkbox"/>	EX
Cytoplasm	Cytoplasm		3	1		<input checked="" type="checkbox"/>	M

3.1 Compartment EX

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

Name Extracellular

3.2 Compartment M

This is a two dimensional compartment with a constant size of one μm^2 , which is surrounded by EX (Extracellular).

Name Membrane

3.3 Compartment Cytoplasm

This is a three dimensional compartment with a constant size of one litre, which is surrounded by M (Membrane).

Name Cytoplasm

4 Species

This model contains 23 species. The boundary condition of one of these species is set to `true` so that this species' amount cannot be changed by any reaction. Section 9 provides further details and the derived rates of change of each species.

Table 3: Properties of each species.

Id	Name	Compartment	Derived Unit	Constant	Boundary Condition
GTPgS_C	GTPgS_C	Cytoplasm	$\mu\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
KCNQsites_M	KCNQsites_M	M	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
PIP2_M	PIP2_M	M	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
GGDPbS_M	GGDPbS_M	M	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
Mg2_C	Mg2_C	Cytoplasm	$\mu\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
ATP_C	ATP_C	Cytoplasm	$\mu\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
oxoM_EX	oxoM_EX	EX	$\mu\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
GDPbS_C	GDPbS_C	Cytoplasm	$\mu\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
GGTPMg_M	GGTPMg_M	M	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
GDP_C	GDP_C	Cytoplasm	$\mu\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
GTP_C	GTP_C	Cytoplasm	$\mu\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
GGDPA1F4Mg_M	GGDPA1F4Mg_M	M	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
ip3_C	IP3_C	Cytoplasm	$\mu\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
A1F4_C	A1F4_C	Cytoplasm	$\mu\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
G_M	G_M	M	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
GGTP_M	GGTP_M	M	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
GGDPA1F4_M	GGDPA1F4_M	M	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
GGTPgS_M	GGTPgS_M	M	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
PI_M	PI_M	M	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
PIP_M	PIP_M	M	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
PIP2xKCNQ_M	PIP2xKCNQ_M	M	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>

Id	Name	Compartment	Derived Unit	Constant	Boundary Condi- tion
GGTPgSMg_M	GGTPgSMg_M	M	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
GGDP_M	GGDP_M	M	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>

5 Parameters

This model contains twelve global parameters.

Table 4: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
MgSat20	MgSat20		0.000	dimensionless	<input type="checkbox"/>
ATPSat300	ATPSat300		0.000	dimensionless	<input type="checkbox"/>
MgSat10	MgSat10		0.000	dimensionless	<input type="checkbox"/>
OxoSat	OxoSat		0.000	dimensionless	<input type="checkbox"/>
Gactive	Gactive		0.000	$\mu\text{m}^{-2} \cdot \text{item}$	<input type="checkbox"/>
ATPSat1000	ATPSat1000		0.000	dimensionless	<input type="checkbox"/>
KMgGTPase	KMgGTPase		10.000	$\mu\text{mol} \cdot \text{l}^{-1}$	<input checked="" type="checkbox"/>
Koxom	Koxom		8.000	$\mu\text{mol} \cdot \text{l}^{-1}$	<input checked="" type="checkbox"/>
I_KCNQ	I_KCNQ		0.000	dimensionless	<input type="checkbox"/>
nHill			1.800	dimensionless	<input checked="" type="checkbox"/>
fGactive	fGactive		0.000	dimensionless	<input type="checkbox"/>
NA_micro	NA(micro)		$6.022 \cdot 10^{17}$	$\text{item} \cdot \mu\text{mol}^{-1}$	<input checked="" type="checkbox"/>

6 Rules

This is an overview of nine rules.

6.1 Rule MgSat20

Rule MgSat20 is an assignment rule for parameter MgSat20:

$$\text{MgSat20} = 1 \cdot \frac{1}{1 + 20 \cdot \frac{1}{[\text{Mg2.C}]}} \quad (1)$$

6.2 Rule ATPSat1000

Rule ATPSat1000 is an assignment rule for parameter ATPSat1000:

$$\text{ATPSat1000} = 1 \cdot \frac{1}{1 + 1000 \cdot \frac{1}{0.5 + [\text{ATP.C}]}} \quad (2)$$

6.3 Rule MgSat10

Rule MgSat10 is an assignment rule for parameter MgSat10:

$$\text{MgSat10} = 0.2 + 0.8 \cdot 1 \cdot \frac{1}{1 + \text{KMgGTPase} \cdot \frac{1}{[\text{Mg2.C}]}} \quad (3)$$

6.4 Rule OxoSat

Rule OxoSat is an assignment rule for parameter OxoSat:

$$\text{OxoSat} = 1 \cdot \frac{1}{1 + \text{Koxom} \cdot \frac{1}{1.0E-5 + [\text{oxoM.EX}]}} \quad (4)$$

6.5 Rule fGactive

Rule fGactive is an assignment rule for parameter fGactive:

$$\text{fGactive} = \frac{[\text{GGTPMg_M}] + [\text{GGTPgSMg_M}] + [\text{GGDPAIF4Mg_M}]}{200} \quad (5)$$

6.6 Rule Gactive

Rule Gactive is an assignment rule for parameter Gactive:

$$\text{Gactive} = [\text{GGTPMg_M}] + [\text{GGTPgSMg_M}] + [\text{GGDPAIF4Mg_M}] \quad (6)$$

Derived unit item · μm⁻²

6.7 Rule ATPSat300

Rule ATPSat300 is an assignment rule for parameter ATPSat300:

$$\text{ATPSat300} = 1 \cdot \frac{1}{1 + 300 \cdot \frac{1}{0.5 + [\text{ATP.C}]}} \quad (7)$$

6.8 Rule I_KCNQ

Rule I_KCNQ is an assignment rule for parameter I_KCNQ:

$$\text{I_KCNQ} = 1.33 \cdot (0.025 \cdot [\text{PIP2xKCNQ_M}])^{\text{nHill}} \quad (8)$$

6.9 Rule GTP_C

Rule GTP_C is an assignment rule for species GTP_C:

$$\text{GTP_C} = 100 - 50 \cdot \exp((0.008333333333333333 \cdot (290 + t))) \quad (9)$$

7 Events

This is an overview of two events. Each event is initiated whenever its trigger condition switches from false to true. A delay function postpones the effects of an event to a later time point. At the time of execution, an event can assign values to species, parameters or compartments if these are not set to constant.

7.1 Event `event_0000001`

Trigger condition

$$t \geq 3 \quad (10)$$

Assignment

$$\text{oxoM_EX} = 10 \quad (11)$$

7.2 Event `event_0000002`

Trigger condition

$$t \geq 8 \quad (12)$$

Assignment

$$\text{oxoM_EX} = 0 \quad (13)$$

8 Reactions

This model contains 15 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

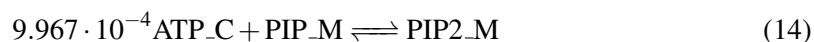
Nº	Id	Name	Reaction Equation	SBO
1	PIP5kinase	PIP5kinase	$9.967 \cdot 10^{-4} \text{ATP_C} + \text{PIP_M} \rightleftharpoons \text{PIP2_M}$	
2	IP3Phosphatase	IP3Phosphatase	$1.666111296 \cdot 10^{-18} \text{ip3_C} \rightleftharpoons \emptyset$	
3	PIP2binding	PIP2binding	$\text{KCNQsites_M} + \text{PIP2_M} \rightleftharpoons \text{PIP2xKCNQ_M}$	
4	MgonGGTPgS	MgonGGTPgS	$\text{GGTPgS_M} \xrightleftharpoons{\text{Mg2_C}} \text{GGTPgSMg_M}$	
5	GGTPase	GGTPase	$\text{GGTPMg_M} \rightleftharpoons \text{GGDP_M}$	
6	GTPon	GTPon	$\text{G_M} \xrightleftharpoons{\text{GTP_C}} \text{GGTP_M}$	
7	GTPgSon	GTPgSon	$9.967 \cdot 10^{-4} \text{GTPgS_C} + \text{G_M} \rightleftharpoons \text{GGTPgS_M}$	
8	GDPbSon	GDPbSon	$\text{G_M} + 9.967 \cdot 10^{-4} \text{GDPbS_C} \rightleftharpoons \text{GGDPbS_M}$	
9	MgGTPon	MgGTPon	$\text{GGTP_M} \xrightleftharpoons{\text{Mg2_C}} \text{GGTPMg_M}$	
10	GDPdissoc	GDPdissoc	$\text{GGDP_M} \xrightleftharpoons{\text{oxoM_EX}} 9.967 \cdot 10^{-4} \text{GDP_C} + \text{G_M}$	
11	PIP2_5Pase	PIP2_5Pase	$\text{PIP2_M} \rightleftharpoons \text{PIP_M}$	
12	PLC	PLC	$\text{PIP2_M} \rightleftharpoons 9.967 \cdot 10^{-4} \text{ip3_C}$	
13	AlF4on	AlF4on	$9.967 \cdot 10^{-4} \text{AlF4_C}$	+
			$\text{GGDP_M} \rightleftharpoons \text{GGDPAIF4_M}$	
14	PI4Kinase	PI4Kinase	$9.967 \cdot 10^{-4} \text{ATP_C} + \text{PI_M} \rightleftharpoons \text{PIP_M}$	
15	MgonGDPAIF4	MgonGDPAIF4	$\text{GGDPAIF4_M} \xrightleftharpoons{\text{Mg2_C}} \text{GGDPAIF4Mg_M}$	

8.1 Reaction PIP5kinase

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

Name PIP5kinase

Reaction equation



Reactants

Table 6: Properties of each reactant.

Id	Name	SBO
ATP_C	ATP_C	
PIP_M	PIP_M	

Product

Table 7: Properties of each product.

Id	Name	SBO
PIP2_M	PIP2_M	

Kinetic Law

Derived unit contains undeclared units

$$v_1 = (0.2 + 0.8 \cdot \text{MgSat20}) \cdot \text{kPIP5Kinase} \cdot [\text{PIP_M}] \cdot \text{ATPSat300} \quad (15)$$

Table 8: Properties of each parameter.

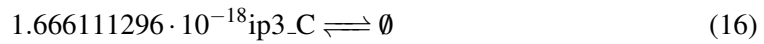
Id	Name	SBO	Value	Unit	Constant
kPIP5Kinase			0.06	s ⁻¹	<input checked="" type="checkbox"/>

8.2 Reaction IP3Phosphatase

This is a reversible reaction of one reactant forming no product.

Name IP3Phosphatase

Reaction equation



Reactant

Table 9: Properties of each reactant.

Id	Name	SBO
ip3_C	IP3_C	

Kinetic Law

Derived unit $\text{s}^{-1} \cdot \text{item}$

$$v_2 = [\text{ip3_C}] \cdot \text{kIP3ase} \cdot \text{vol}(\text{Cytoplasm}) \cdot \text{NA_micro} \quad (17)$$

Table 10: Properties of each parameter.

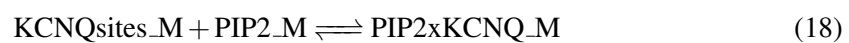
Id	Name	SBO	Value	Unit	Constant
kIP3ase			0.3	s^{-1}	<input checked="" type="checkbox"/>

8.3 Reaction PIP2binding

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

Name PIP2binding

Reaction equation



Reactants

Table 11: Properties of each reactant.

Id	Name	SBO
KCNQsites_M	KCNQsites_M	
PIP2_M	PIP2_M	

Product

Table 12: Properties of each product.

Id	Name	SBO
PIP2xKCNQ_M	PIP2xKCNQ_M	

Kinetic Law

Derived unit $\text{s}^{-1} \cdot \text{item}$

$$v_3 = (\text{kPIP2on} \cdot [\text{KCNQsites_M}] \cdot [\text{PIP2_M}] + ((\text{kPIP2off} \cdot [\text{PIP2xKCNQ_M}]))) \cdot \text{area}(\text{M}) \quad (19)$$

Table 13: Properties of each parameter.

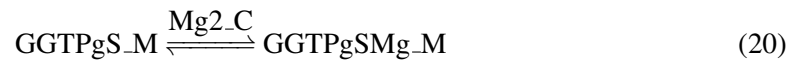
Id	Name	SBO	Value	Unit	Constant
kPIP2on			$2.5 \cdot 10^{-4}$	$\text{item}^{-1} \cdot \mu\text{m}^2 \cdot \text{s}^{-1}$	<input checked="" type="checkbox"/>
kPIP2off			0.250	s^{-1}	<input checked="" type="checkbox"/>

8.4 Reaction MgonGGTPgS

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

Name MgonGGTPgS

Reaction equation



Reactant

Table 14: Properties of each reactant.

Id	Name	SBO
GGTPgS_M	GGTPgS_M	

Modifier

Table 15: Properties of each modifier.

Id	Name	SBO
Mg2_C	Mg2_C	

Product

Table 16: Properties of each product.

Id	Name	SBO
GGTPgSMg_M	GGTPgSMg_M	

Kinetic Law

Derived unit $\text{s}^{-1} \cdot \text{item}$

$$v_4 = \text{kMg2onGTPgS} \cdot [\text{GGTPgS_M}] \cdot [\text{Mg2_C}] \cdot \text{area (M)} \quad (21)$$

Table 17: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
kMg2onGTPgS			0.002	$1 \cdot \mu\text{mol}^{-1} \cdot \text{s}^{-1}$	<input checked="" type="checkbox"/>

8.5 Reaction GGTPase

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

Name GGTPase

Reaction equation



Reactant

Table 18: Properties of each reactant.

Id	Name	SBO
GGTPMg_M	GGTPMg_M	

Product

Table 19: Properties of each product.

Id	Name	SBO
GGDP_M	GGDP_M	

Id	Name	SBO
----	------	-----

Kinetic Law

Derived unit $\text{s}^{-1} \cdot \text{item}$

$$v_5 = k\text{GGTPase} \cdot \text{MgSat10} \cdot [\text{GGTPMg_M}] \cdot \text{area}(\text{M}) \quad (23)$$

Table 20: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
kGGTPase	kGGTPase		1.8	s^{-1}	<input checked="" type="checkbox"/>

8.6 Reaction GTPon

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

Name GTPon

Reaction equation



Reactant

Table 21: Properties of each reactant.

Id	Name	SBO
G_M	G_M	

Modifier

Table 22: Properties of each modifier.

Id	Name	SBO
GTP_C	GTP_C	

Product

Table 23: Properties of each product.

Id	Name	SBO
GGTP_M	GGTP_M	

Kinetic Law

Derived unit $\text{s}^{-1} \cdot \text{item}$

$$v_6 = (\text{kGTPon} \cdot [\text{G_M}] \cdot [\text{GTP_C}] + ((\text{kGTPoff} \cdot [\text{GGTP_M}]))) \cdot \text{area}(\text{M}) \quad (25)$$

Table 24: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
kGTPon			0.45	$1 \cdot \mu\text{mol}^{-1} \cdot \text{s}^{-1}$	<input checked="" type="checkbox"/>
kGTPoff			0.08	s^{-1}	<input checked="" type="checkbox"/>

8.7 Reaction GTPgSon

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

Name GTPgSon

Reaction equation



Reactants

Table 25: Properties of each reactant.

Id	Name	SBO
GTPgS_C	GTPgS_C	
G_M	G_M	

Product

Table 26: Properties of each product.

Id	Name	SBO
GGTPgS_M	GGTPgS_M	

Kinetic Law

Derived unit $\text{s}^{-1} \cdot \text{item}$

$$v_7 = (\text{kGTPgSon} \cdot [\text{G_M}] \cdot [\text{GTPgS_C}] + ((\text{kGTPgSoff} \cdot [\text{GGTPgS_M}]))) \cdot \text{area}(\text{M}) \quad (27)$$

Table 27: Properties of each parameter.

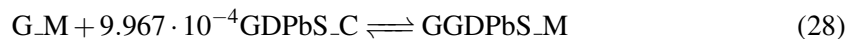
Id	Name	SBO	Value	Unit	Constant
kGTPgSon			0.006	$1 \cdot \mu\text{mol}^{-1} \cdot \text{s}^{-1}$	<input checked="" type="checkbox"/>
kGTPgSoff			0.005	s^{-1}	<input checked="" type="checkbox"/>

8.8 Reaction GDPbSon

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

Name GDPbSon

Reaction equation



Reactants

Table 28: Properties of each reactant.

Id	Name	SBO
G_M	G_M	
GDPbS_C	GDPbS_C	

Product

Table 29: Properties of each product.

Id	Name	SBO
GGDPbS_M	GGDPbS_M	

Kinetic Law

Derived unit $\text{s}^{-1} \cdot \text{item}$

$$v_8 = (kGDPbSon \cdot [G_M] \cdot [GDPbS_C] + ((kGGDPbSoff \cdot (1 + 20 \cdot OxoSat) \cdot [GGDPbS_M]))) \cdot area(M) \quad (29)$$

Table 30: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
kGDPbSon			0.28	$l \cdot \mu\text{mol}^{-1} \cdot s^{-1}$	<input checked="" type="checkbox"/>
kGGDPbSoff			0.10	s^{-1}	<input checked="" type="checkbox"/>

8.9 Reaction MgGTPon

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

Name MgGTPon

Reaction equation



Reactant

Table 31: Properties of each reactant.

Id	Name	SBO
GGTP_M	GGTP_M	

Modifier

Table 32: Properties of each modifier.

Id	Name	SBO
Mg2_C	Mg2_C	

Product

Table 33: Properties of each product.

Id	Name	SBO
GGTPMg_M	GGTPMg_M	

Kinetic Law

Derived unit $\text{s}^{-1} \cdot \text{item}$

$$v_9 = \text{kMg2onGTP} \cdot [\text{GGTP_M}] \cdot [\text{Mg2_C}] \cdot \text{area}(\text{M}) \quad (31)$$

Table 34: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
kMg2onGTP			0.003	$1 \cdot \mu\text{mol}^{-1} \cdot \text{s}^{-1}$	<input checked="" type="checkbox"/>

8.10 Reaction GDPdissoc

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

Name GDPdissoc

Reaction equation



Reactant

Table 35: Properties of each reactant.

Id	Name	SBO
GGDP_M	GGDP_M	

Modifier

Table 36: Properties of each modifier.

Id	Name	SBO
oxoM_EX	oxoM_EX	

Products

Table 37: Properties of each product.

Id	Name	SBO
GDP_C	GDP_C	

Id	Name	SBO
G_M	G_M	

Kinetic Law

Derived unit $\text{s}^{-1} \cdot \text{item}$

$$v_{10} = (\text{kGDPoff} \cdot [\text{GGDP_M}] \cdot (\text{OxoSat} + \text{TonicAct}) + ((\text{kGDPon} \cdot [\text{GDP_C}] \cdot [\text{G_M}]))) \cdot \text{area (M)} \quad (33)$$

Table 38: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
kGDPoff			0.500	s^{-1}	<input checked="" type="checkbox"/>
TonicAct			0.002	dimensionless	<input checked="" type="checkbox"/>
kGDPon			0.003	$1 \cdot \mu\text{mol}^{-1} \cdot \text{s}^{-1}$	<input checked="" type="checkbox"/>

8.11 Reaction PIP2_5Pase

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

Name PIP2_5Pase

Reaction equation



Reactant

Table 39: Properties of each reactant.

Id	Name	SBO
PIP2_M	PIP2_M	

Product

Table 40: Properties of each product.

Id	Name	SBO
PIP_M	PIP_M	

Kinetic Law

Derived unit $\text{s}^{-1} \cdot \text{item}$

$$v_{11} = \text{MgSat20} \cdot \text{kPIP2Pase} \cdot [\text{PIP2_M}] \cdot \text{area}(\text{M}) \quad (35)$$

Table 41: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
kPIP2Pase			0.005	s^{-1}	<input checked="" type="checkbox"/>

8.12 Reaction PLC

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

Name PLC

Reaction equation



Reactant

Table 42: Properties of each reactant.

Id	Name	SBO
PIP2_M	PIP2_M	

Product

Table 43: Properties of each product.

Id	Name	SBO
ip3_C	IP3_C	

Kinetic Law

Derived unit $\text{s}^{-1} \cdot \text{item}$

$$v_{12} = \text{kPLC} \cdot (\text{fGactive} + \text{PLCspont}) \cdot [\text{PIP2_M}] \cdot \text{area}(\text{M}) \quad (37)$$

Table 44: Properties of each parameter.

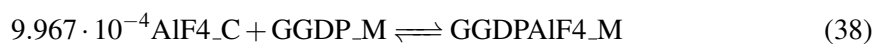
Id	Name	SBO	Value	Unit	Constant
kPLC			4.800	s ⁻¹	<input checked="" type="checkbox"/>
PLCspont			7.5 · 10 ⁻⁴	dimensionless	<input checked="" type="checkbox"/>

8.13 Reaction A1F4on

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

Name A1F4on

Reaction equation



Reactants

Table 45: Properties of each reactant.

Id	Name	SBO
A1F4_C	A1F4_C	
GGDP_M	GGDP_M	

Product

Table 46: Properties of each product.

Id	Name	SBO
GGDPA1F4_M	GGDPA1F4_M	

Kinetic Law

Derived unit s⁻¹ · item

$$v_{13} = (k\text{A1F4on} \cdot [\text{A1F4_C}] \cdot [\text{GGDP_M}] + ((k\text{A1F4off} \cdot [\text{GGDPA1F4_M}]))) \cdot \text{area}(\text{M}) \quad (39)$$

Table 47: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
kA1F4on			7 · 10 ⁻⁶	1 · μmol ⁻¹ · s ⁻¹	<input checked="" type="checkbox"/>
kA1F4off			0.005	s ⁻¹	<input checked="" type="checkbox"/>

8.14 Reaction PI4Kinase

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

Name PI4Kinase

Reaction equation



Reactants

Table 48: Properties of each reactant.

Id	Name	SBO
ATP_C	ATP_C	
PI_M	PI_M	

Product

Table 49: Properties of each product.

Id	Name	SBO
PIP_M	PIP_M	

Kinetic Law

Derived unit contains undeclared units

$$v_{14} = (0.2 + 0.8 \cdot \text{MgSat20}) \cdot \text{kPI4Kinase} \cdot [\text{PI_M}] \cdot \text{ATPSat1000} \cdot \text{area (M)} \quad (41)$$

Table 50: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
kPI4Kinase			10^{-4}	s^{-1}	<input checked="" type="checkbox"/>

8.15 Reaction MgonGDPAIF4

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

Name MgonGDPAIF4

Reaction equation



Reactant

Table 51: Properties of each reactant.

Id	Name	SBO
GGDPA1F4_M	GGDPAIF4_M	

Modifier

Table 52: Properties of each modifier.

Id	Name	SBO
Mg2_C	Mg2_C	

Product

Table 53: Properties of each product.

Id	Name	SBO
GGDPA1F4Mg_M	GGDPAIF4Mg_M	

Kinetic Law

Derived unit $\text{s}^{-1} \cdot \text{item}$

$$v_{15} = \text{kMgonAlF4} \cdot [\text{GGDPAIF4_M}] \cdot [\text{Mg2_C}] \cdot \text{area (M)} \quad (43)$$

Table 54: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
kMgonAlF4			0.002	$1 \cdot \mu\text{mol}^{-1} \cdot \text{s}^{-1}$	<input checked="" type="checkbox"/>

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 `GTPgS_C`

Name `GTPgS_C`

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

This species takes part in one reaction (as a reactant in `GTPgSon`).

$$\frac{d}{dt} \text{GTPgS_C} = -9.967 \cdot 10^{-4} v_7 \quad (44)$$

9.2 Species `KCNQsites_M`

Name `KCNQsites_M`

Initial concentration $6.7 \text{ item} \cdot \mu\text{m}^{-2}$

This species takes part in one reaction (as a reactant in `PIP2binding`).

$$\frac{d}{dt} \text{KCNQsites_M} = -v_3 \quad (45)$$

9.3 Species `PIP2_M`

Name `PIP2_M`

Initial concentration $5000 \text{ item} \cdot \mu\text{m}^{-2}$

This species takes part in four reactions (as a reactant in `PIP2binding`, `PIP2_5Pase`, `PLC` and as a product in `PIP5kinase`).

$$\frac{d}{dt} \text{PIP2_M} = v_1 - v_3 - v_{11} - v_{12} \quad (46)$$

9.4 Species [GGDPbS_M](#)

Name GGDPbS_M

Initial concentration 0 item · μm⁻²

This species takes part in one reaction (as a product in [GDPbSon](#)).

$$\frac{d}{dt} \text{GGDPbS_M} = v_8 \quad (47)$$

9.5 Species [Mg2_C](#)

Name Mg2_C

Initial concentration 2100 μmol · l⁻¹

This species takes part in three reactions (as a modifier in [MgonGGTPgS](#), [MgGTPon](#), [MgonGDPA1F4](#)), which do not influence its rate of change because this species is on the boundary of the reaction system:

$$\frac{d}{dt} \text{Mg2_C} = 0 \quad (48)$$

9.6 Species [ATP_C](#)

Name ATP_C

Initial concentration 3000 μmol · l⁻¹

This species takes part in two reactions (as a reactant in [PIP5kinase](#), [PI4Kinase](#)).

$$\frac{d}{dt} \text{ATP_C} = -9.967 \cdot 10^{-4} v_1 - 9.967 \cdot 10^{-4} v_{14} \quad (49)$$

9.7 Species [oxoM_EX](#)

Name oxoM_EX

Initial concentration 0 μmol · l⁻¹

Involved in events [event_0000001](#), [event_0000002](#)

This species takes part in one reaction (as a modifier in [GDPdissoc](#)).

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

Furthermore, two events influence this species' rate of change.

9.8 Species [GDPbS_C](#)

Name GDPbS_C

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

This species takes part in one reaction (as a reactant in [GDPbSon](#)).

$$\frac{d}{dt}\text{GDPbS_C} = -9.967 \cdot 10^{-4} v_8 \quad (51)$$

9.9 Species [GGTPMg_M](#)

Name GGTPMg_M

Initial concentration 0.1 $\text{item} \cdot \mu\text{m}^{-2}$

This species takes part in two reactions (as a reactant in [GGTPase](#) and as a product in [MgGTPon](#)).

$$\frac{d}{dt}\text{GGTPMg_M} = v_9 - v_5 \quad (52)$$

9.10 Species [GDP_C](#)

Name GDP_C

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

This species takes part in one reaction (as a product in [GDPdissoc](#)).

$$\frac{d}{dt}\text{GDP_C} = 9.967 \cdot 10^{-4} v_{10} \quad (53)$$

9.11 Species [GTP_C](#)

Name GTP_C

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

Involved in rule [GTP_C](#)

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

9.12 Species [GGDPAIF4Mg_M](#)

Name GGDPAIF4Mg_M

Initial concentration 0 $\text{item} \cdot \mu\text{m}^{-2}$

This species takes part in one reaction (as a product in [MgonGDPAlF4](#)).

$$\frac{d}{dt}\text{GGDPAIF4Mg_M} = v_{15} \quad (54)$$

9.13 Species ip3_C

Name IP3_C

Initial concentration $0.48 \mu\text{mol} \cdot \text{l}^{-1}$

This species takes part in two reactions (as a reactant in [IP3Phosphatase](#) and as a product in [PLC](#)).

$$\frac{d}{dt}\text{ip3_C} = 9.967 \cdot 10^{-4} v_{12} - 1.666111296 \cdot 10^{-18} v_2 \quad (55)$$

9.14 Species AIF4_C

Name AIF4_C

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

This species takes part in one reaction (as a reactant in [AlF4on](#)).

$$\frac{d}{dt}\text{AIF4_C} = -9.967 \cdot 10^{-4} v_{13} \quad (56)$$

9.15 Species G_M

Name G_M

Initial concentration $0.065 \text{ item} \cdot \mu\text{m}^{-2}$

This species takes part in four reactions (as a reactant in [GTPon](#), [GTPgSon](#), [GDPbSon](#) and as a product in [GDPdissoc](#)).

$$\frac{d}{dt}\text{G_M} = v_{10} - v_6 - v_7 - v_8 \quad (57)$$

9.16 Species GGTP_M

Name GGTP_M

Initial concentration $0.048 \text{ item} \cdot \mu\text{m}^{-2}$

This species takes part in two reactions (as a reactant in [MgGTPon](#) and as a product in [GTPon](#)).

$$\frac{d}{dt}\text{GGTP_M} = v_6 - v_9 \quad (58)$$

9.17 Species GGDPAIF4_M

Name GGDPAIF4_M

Initial concentration 0 item · μm⁻²

This species takes part in two reactions (as a reactant in [MgonGGDPAIF4](#) and as a product in [AlF4on](#)).

$$\frac{d}{dt}\text{GGDPAIF4_M} = v_{13} - v_{15} \quad (59)$$

9.18 Species GGTPgS_M

Name GGTPgS_M

Initial concentration 0 item · μm⁻²

This species takes part in two reactions (as a reactant in [MgonGGTPgS](#) and as a product in [GTPgSon](#)).

$$\frac{d}{dt}\text{GGTPgS_M} = v_7 - v_4 \quad (60)$$

9.19 Species PI_M

Name PI_M

Initial concentration 200000 item · μm⁻²

This species takes part in one reaction (as a reactant in [PI4Kinase](#)).

$$\frac{d}{dt}\text{PI_M} = -v_{14} \quad (61)$$

9.20 Species PIP_M

Name PIP_M

Initial concentration 1150 item · μm⁻²

This species takes part in three reactions (as a reactant in [PIP5kinase](#) and as a product in [PIP2.5Pase](#), [PI4Kinase](#)).

$$\frac{d}{dt}\text{PIP_M} = v_{11} + v_{14} - v_1 \quad (62)$$

9.21 Species PIP2xKCNQ_M

Name PIP2xKCNQ_M

Initial concentration 33.3 item · μm⁻²

This species takes part in one reaction (as a product in [PIP2binding](#)).

$$\frac{d}{dt} \text{PIP2xKCNQ_M} = v_3 \quad (63)$$

9.22 Species GGTPgSMg_M

Name GGTPgSMg_M

Initial concentration 0 item · μm⁻²

This species takes part in one reaction (as a product in [MgonGGTPgS](#)).

$$\frac{d}{dt} \text{GGTPgSMg_M} = v_4 \quad (64)$$

9.23 Species GGDP_M

Name GGDP_M

Initial concentration 200 item · μm⁻²

This species takes part in three reactions (as a reactant in [GDPdissoc](#), [AlF4on](#) and as a product in [GGTPase](#)).

$$\frac{d}{dt} \text{GGDP_M} = v_5 - v_{10} - v_{13} \quad (65)$$

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