

SBML Model Report

Model name: “Eungdamrong2007_Ras_Activation”



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 January 29th 2008 at 10:13 a.m. and last time modified at March 20th 2014 at 5:03 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	7
species types	0	species	46
events	0	constraints	0
reactions	43	function definitions	0
global parameters	26	unit definitions	21
rules	12	initial assignments	0

Model Notes

The model reproduces the time profiles of Golgi Ras-GTP and plasma membrane Ras-GTP, subjected to a palmitoylation rate of 0.00015849 second inverse. This is depicted in Fig 5a and 5b for various palmitoylation rates, however the value used in this model is not present in the figure in the paper but corresponds to Fig S2 of the supplement. Model successfully reproduced using MathSBML. Please note that the units of volumetric species in this model are

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

molecules/micrometer cubed, to convert this to microMolar as given in the paper, multiply the simulation result by 1/602.

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2 Unit Definitions

This is an overview of 23 unit definitions of which two are predefined by SBML and not mentioned in the model.

2.1 Unit substance

Definition item

2.2 Unit volume

Definition μm^3

2.3 Unit area

Definition μm^2

2.4 Unit molecules

Definition item

2.5 Unit `umol_um3_litre_1`

Definition 10^{-21} mol

2.6 Unit `um2`

Definition μm^2

2.7 Unit `uM_um3_molecules_1`

Definition $10^{-21} \text{ dimensionless} \cdot \text{item}^{-1} \cdot \text{mol}$

2.8 Unit `molecules_um_2_s_1`

Definition $10^{12} \text{ dimensionless} \cdot \text{item} \cdot \text{m}^{-2} \cdot \text{s}^{-1}$

2.9 Unit `pA_um_2`

Definition $\text{dimensionless} \cdot \text{A} \cdot \text{m}^{-2}$

2.10 Unit `s_1`

Definition s^{-1}

2.11 Unit `s`

Definition s

2.12 Unit `molecules_um_2`

Definition $10^{12} \text{ dimensionless} \cdot \text{item} \cdot \text{m}^{-2}$

2.13 Unit `uM_s_1`

Definition $0.0010 \text{ dimensionless} \cdot \text{m}^{-3} \cdot \text{mol} \cdot \text{s}^{-1}$

2.14 Unit `uM_1_s_1`

Definition $1000 \text{ dimensionless} \cdot \text{m}^3 \cdot \text{mol}^{-1} \cdot \text{s}^{-1}$

2.15 Unit `uM`

Definition $0.0010 \text{ dimensionless} \cdot \text{m}^{-3} \cdot \text{mol}$

2.16 Unit `molecules_um_2_uM_1_s_1`

Definition $10^{15} \text{ dimensionless} \cdot \text{item} \cdot \text{m} \cdot \text{mol}^{-1} \cdot \text{s}^{-1}$

2.17 Unit `um2_molecules_1_s_1`

Definition $10^{-12} \text{ dimensionless} \cdot \text{item}^{-1} \cdot \text{m}^2 \cdot \text{s}^{-1}$

2.18 Unit `uM_um_s_1`

Definition $10^{-9} \text{ dimensionless} \cdot \text{m}^{-2} \cdot \text{mol} \cdot \text{s}^{-1}$

2.19 Unit `_one_0E33_item_2_m2_mol_s_1`

Definition $10^{-33} \text{ dimensionless} \cdot \text{item}^{-2} \cdot \text{m}^2 \cdot \text{mol} \cdot \text{s}^{-1}$

2.20 Unit `_one_0E30_item_2_m5_s_1`

Definition $10^{-30} \text{ dimensionless} \cdot \text{item}^{-2} \cdot \text{m}^5 \cdot \text{s}^{-1}$

2.21 Unit `um3_molecules_1_s_1`

Definition $10^{-18} \text{ dimensionless} \cdot \text{item}^{-1} \cdot \text{m}^3 \cdot \text{s}^{-1}$

2.22 Unit `length`

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

Definition m

2.23 Unit `time`

Notes Second is the predefined SBML unit for time.

Definition s

3 Compartments

This model contains seven compartments.

Table 2: Properties of all compartments.

Id	Name	SBO	Spatial Dimensions	Size	Unit	Constant	Outside
EC	EC		3	1	μm^3	<input checked="" type="checkbox"/>	
cyt	cyt		3	0.976	μm^3	<input checked="" type="checkbox"/>	PM
er	er		3	0.012	μm^3	<input checked="" type="checkbox"/>	erMembrane
Golgi	Golgi		3	0.012	μm^3	<input checked="" type="checkbox"/>	GM
PM	PM		2	0.6	μm^2	<input checked="" type="checkbox"/>	EC
erMembrane	erMembrane		2	0.0456	μm^2	<input checked="" type="checkbox"/>	cyt
GM	GM		2	0.048	μm^2	<input checked="" type="checkbox"/>	cyt

3.1 Compartment `EC`

This is a three dimensional compartment with a constant size of one μm^3 .

Name EC

3.2 Compartment *cyt*

This is a three dimensional compartment with a constant size of $0.976 \mu\text{m}^3$, which is surrounded by PM (PM).

Name *cyt*

3.3 Compartment *er*

This is a three dimensional compartment with a constant size of $0.012 \mu\text{m}^3$, which is surrounded by erMembrane (erMembrane).

Name *er*

3.4 Compartment *Golgi*

This is a three dimensional compartment with a constant size of $0.012 \mu\text{m}^3$, which is surrounded by GM (GM).

Name *Golgi*

3.5 Compartment *PM*

This is a two dimensional compartment with a constant size of $0.6 \mu\text{m}^2$, which is surrounded by EC (EC).

Name *PM*

3.6 Compartment *erMembrane*

This is a two dimensional compartment with a constant size of $0.0456 \mu\text{m}^2$, which is surrounded by *cyt* (*cyt*).

Name *erMembrane*

3.7 Compartment *GM*

This is a two dimensional compartment with a constant size of $0.048 \mu\text{m}^2$, which is surrounded by *cyt* (*cyt*).

Name *GM*

4 Species

This model contains 46 species. The boundary condition of three 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
RasGTP_Golgi_GM		GM	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
EGF_EC		EC	$\text{item} \cdot \mu\text{m}^{-3}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
CAPRI_cyt		cyt	$\text{item} \cdot \mu\text{m}^{-3}$	<input type="checkbox"/>	<input type="checkbox"/>
serca		erMembrane	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
PIP_PM		PM	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
PI_PM		PM	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
Shc_PM		PM	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
CaCAPRI_PM_PM		PM	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
RactCa		erMembrane	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
Shc_star_PM		PM	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
EGFR_PM		PM	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
PLC_act_PM		PM	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
RasGTP_pal_cyt		cyt	$\text{item} \cdot \mu\text{m}^{-3}$	<input type="checkbox"/>	<input type="checkbox"/>
PLC_PM		PM	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
PIP2_PM		PM	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
Activated_EGFR_PM		PM	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
Ca		cyt	$\text{item} \cdot \mu\text{m}^{-3}$	<input type="checkbox"/>	<input type="checkbox"/>
Ract		erMembrane	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
Rinh		erMembrane	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
RinhCa		erMembrane	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
IP3		cyt	$\text{item} \cdot \mu\text{m}^{-3}$	<input type="checkbox"/>	<input type="checkbox"/>

Id	Name	Compartment	Derived Unit	Constant	Boundary Condi- tion
RasGDP_Golgi_GM		GM	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
Ca_RasGRP_GM_GM		GM	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
DAG_GM_GM		GM	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
RasGRP_DAG_GM		GM	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
CaCAPRI_cyt		cyt	$\text{item} \cdot \mu\text{m}^{-3}$	<input type="checkbox"/>	<input type="checkbox"/>
DAG_PM		PM	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
RasGTP_depall_cyt		cyt	$\text{item} \cdot \mu\text{m}^{-3}$	<input type="checkbox"/>	<input type="checkbox"/>
RasGDP_depall_cyt		cyt	$\text{item} \cdot \mu\text{m}^{-3}$	<input type="checkbox"/>	<input type="checkbox"/>
RasGDP_pall_cyt		cyt	$\text{item} \cdot \mu\text{m}^{-3}$	<input type="checkbox"/>	<input type="checkbox"/>
Ca_PLCe_cyt		cyt	$\text{item} \cdot \mu\text{m}^{-3}$	<input type="checkbox"/>	<input type="checkbox"/>
Ras_CaPLCe_GM		GM	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
PIP2_GM_GM		GM	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
ER_erMembrane		erMembrane	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
Ca_ER		er	$\text{item} \cdot \mu\text{m}^{-3}$	<input type="checkbox"/>	<input type="checkbox"/>
Sos_cyt		cyt	$\text{item} \cdot \mu\text{m}^{-3}$	<input type="checkbox"/>	<input type="checkbox"/>
Grb2_cyt		cyt	$\text{item} \cdot \mu\text{m}^{-3}$	<input type="checkbox"/>	<input type="checkbox"/>
PLCe_cyt		cyt	$\text{item} \cdot \mu\text{m}^{-3}$	<input type="checkbox"/>	<input type="checkbox"/>
buffer_cyt		cyt	$\text{item} \cdot \mu\text{m}^{-3}$	<input type="checkbox"/>	<input type="checkbox"/>
ca_buffer_cyt		cyt	$\text{item} \cdot \mu\text{m}^{-3}$	<input type="checkbox"/>	<input type="checkbox"/>
SosGrb2_cyt		cyt	$\text{item} \cdot \mu\text{m}^{-3}$	<input type="checkbox"/>	<input type="checkbox"/>
SGS_PM		PM	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
RasGTP_PM		PM	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
RasGDP_PM		PM	$\text{item} \cdot \mu\text{m}^{-2}$	<input type="checkbox"/>	<input type="checkbox"/>
RasGRP_cyt		cyt	$\text{item} \cdot \mu\text{m}^{-3}$	<input type="checkbox"/>	<input type="checkbox"/>
CaRasGRP1_cyt		cyt	$\text{item} \cdot \mu\text{m}^{-3}$	<input type="checkbox"/>	<input type="checkbox"/>

5 Parameters

This model contains 26 global parameters.

Table 4: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
KMOLE			0.002	10^{-21} dimensionless · item ⁻¹ · mol	✓
kStimSynPIP2- _PIP2- _synthesis			0.010	s ⁻¹	✓
tauPIP2syn- _PIP2- _synthesis			0.050	s	✓
PIP2syndecay- _PIP2- _synthesis			10.000	s	✓
PIP2_basal- _PIP2- _synthesis			1072.000	10^{12} dimensionless · item · m ⁻²	✓
kBasalSynPIP2- _PIP2- _synthesis			0.045	s ⁻¹	✓
Rate- _PIP2Synbasal- _PIP2- _synthesis			0.000	s ⁻¹	☐
Rate- _PIP2SynStim- _PIP2- _synthesis			0.000	s ⁻¹	☐
Vmax_Shc- _phosphorylation			0.000	10^{12} dimensionless · item · m ⁻² · s ⁻¹	☐
Vmax_RasGRP- _DAG_GEF			0.000	10^{12} dimensionless · item · m ⁻² · s ⁻¹	☐
Vmax_CAPRI- _GAP			0.000	10^{12} dimensionless · item · m ⁻² · s ⁻¹	☐
Vmax- _CaRasGRP- _act_RasGM			0.000	10^{12} dimensionless · item · m ⁻² · s ⁻¹	☐
PIP_basal- _PIP- _synthesis			2857.000	10^{12} dimensionless · item · m ⁻²	✓

Id	Name	SBO	Value	Unit	Constant
kBasalSynPIP- _PIP- _synthesis			0.002	s^{-1}	<input checked="" type="checkbox"/>
kStimSynPIP- _PIP- _synthesis			0.010	s^{-1}	<input checked="" type="checkbox"/>
tauPIPsyn- _PIP- _synthesis			0.050	s	<input checked="" type="checkbox"/>
PIPsyndecay- _PIP- _synthesis			10.000	s	<input checked="" type="checkbox"/>
Ratebasal- _PIPsyn_PIP- _synthesis			0.000	s^{-1}	<input type="checkbox"/>
Ratestim- _PIPsyn_PIP- _synthesis			0.000	s^{-1}	<input type="checkbox"/>
Kon- _reaction2			2.100	$1000 \text{ dimensionless} \cdot \text{m}^3 \cdot \text{mol}^{-1} \cdot \text{s}^{-1}$	<input checked="" type="checkbox"/>
dinh- _reaction2			0.110	$0.0010 \text{ dimensionless} \cdot \text{m}^{-3} \cdot \text{mol}$	<input checked="" type="checkbox"/>
Koff- _reaction2			0.000	s^{-1}	<input type="checkbox"/>
Vmax_EGF- _act_PLCGamma			0.000	$10^{12} \text{ dimensionless} \cdot \text{item} \cdot \text{m}^{-2} \cdot \text{s}^{-1}$	<input type="checkbox"/>
dact_Ca- _binds_IP3R			0.200	s^{-1}	<input checked="" type="checkbox"/>
Kr_Ca.binds- _IP3R			0.000	s^{-1}	<input type="checkbox"/>
Vmax_Sos- _act_RasPM			0.000	$10^{12} \text{ dimensionless} \cdot \text{item} \cdot \text{m}^{-2} \cdot \text{s}^{-1}$	<input type="checkbox"/>

6 Rules

This is an overview of twelve rules.

6.1 Rule `Rate_PIP2Synbasal_PIP2_synthesis`

Rule `Rate_PIP2Synbasal_PIP2_synthesis` is an assignment rule for parameter `Rate_PIP2Synbasal_PIP2_synthesis`:

$$\begin{aligned} & \text{Rate_PIP2Synbasal_PIP2_synthesis} & (1) \\ = & \begin{cases} 0.581 \cdot k\text{BasalSynPIP2_PIP2_synthesis} \cdot \left(-1 + \exp \left((\text{PIP2_basal_PIP2_synthesis} + ([\text{PIP2_PM}])) \cdot \frac{1}{\text{PIP2_basal_PIP2_synthesis}} \right) \right) \\ 0 \end{cases} \end{aligned}$$

6.2 Rule `Rate_PIP2SynStim_PIP2_synthesis`

Rule `Rate_PIP2SynStim_PIP2_synthesis` is an assignment rule for parameter `Rate_PIP2SynStim_PIP2_synthesis`:

$$\begin{aligned} & \text{Rate_PIP2SynStim_PIP2_synthesis} & (2) \\ = & \begin{cases} k\text{StimSynPIP2_PIP2_synthesis} \cdot \exp \left(\left((t + (\text{tauPIP2syn_PIP2_synthesis})) \cdot \frac{1}{\text{PIP2syndecay_PIP2_synthesis}} \right) \right) \\ 0 \end{cases} \end{aligned}$$

6.3 Rule `Vmax_Shc_phosphorylation`

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

$$\text{Vmax_Shc_phosphorylation} = 0.2 \cdot [\text{Activated_EGFR_PM}] \quad (3)$$

6.4 Rule `Vmax_RasGRP_DAG_GEF`

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

$$\text{Vmax_RasGRP_DAG_GEF} = 0.05 \cdot [\text{RasGRP_DAG_GM}] \quad (4)$$

6.5 Rule `Vmax_CAPRI_GAP`

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

$$\text{Vmax_CAPRI_GAP} = 10 \cdot [\text{CaCAPRI_PM_PM}] \quad (5)$$

6.6 Rule `Vmax_CaRasGRP_act_RasGM`

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

$$\text{Vmax_CaRasGRP_act_RasGM} = 0.01 \cdot [\text{Ca_RasGRP_GM_GM}] \quad (6)$$

6.7 Rule `Ratebasal_PIPsyn_PIP_synthesis`

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

$$\begin{aligned} & \text{Ratebasal_PIPsyn_PIP_synthesis} & (7) \\ & = \begin{cases} 0.581 \cdot \text{kBasalSynPIP_PIP_synthesis} \cdot \left(-1 + \exp \left((\text{PIP_basal_PIP_synthesis} + ([\text{PIP_PM}])) \cdot \frac{1}{\text{PIP_basal_PIP_synthesis}} \right) \right) \\ 0 \end{cases} \end{aligned}$$

6.8 Rule `Rateestim_PIPsyn_PIP_synthesis`

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

$$\begin{aligned} & \text{Rateestim_PIPsyn_PIP_synthesis} & (8) \\ & = \begin{cases} \text{kStimSynPIP_PIP_synthesis} \cdot \exp \left(\left((t + (\text{tauPIPsyn_PIP_synthesis})) \cdot \frac{1}{\text{PIPsyndecay_PIP_synthesis}} \right) \right) & \text{if } t > \text{tauPIPsyn_PIP_synthesis} \\ 0 & \text{otherwise} \end{cases} \end{aligned}$$

6.9 Rule `Koff_reaction2`

Rule `Koff_reaction2` is an assignment rule for parameter `Koff_reaction2`:

$$\text{Koff_reaction2} = \text{dinh_reaction2} \cdot \text{Kon_reaction2} \quad (9)$$

Derived unit s^{-1}

6.10 Rule `Vmax_EGF_act_PLCgamma`

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

$$\text{Vmax_EGF_act_PLCgamma} = 0.3 \cdot [\text{Activated_EGFR_PM}] \quad (10)$$

6.11 Rule `Kr_Ca_binds_IP3R`

Rule `Kr_Ca_binds_IP3R` is an assignment rule for parameter `Kr_Ca_binds_IP3R`:

$$\text{Kr_Ca_binds_IP3R} = 1000 \cdot \text{dact_Ca_binds_IP3R} \quad (11)$$

6.12 Rule `Vmax_Sos_act_RasPM`

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

$$\text{Vmax_Sos_act_RasPM} = 0.02 \cdot [\text{SGS_PM}] \quad (12)$$

7 Reactions

This model contains 43 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	PIP2_synthesis	PIP2 synthesis	$\text{PIP_PM} \rightleftharpoons \text{PIP2_PM}$	
2	Shc- _phosphorylation	Shc phosphorylation	$\text{Shc_PM} \xrightleftharpoons{\text{Activated_EGFR_PM}} \text{Shc_star_PM}$	
3	Ca_bind_CAPRI	Ca bind CAPRI	$\text{Ca} + \text{CAPRI_cyt} \rightleftharpoons \text{CaCAPRI_cyt}$	
4	IP3_degradation	IP3 degradation	$\text{IP3} \rightleftharpoons \emptyset$	
5	RasGTP_depall- _translocate	RasGTP depal translocate	$\text{RasGTP_depall_cyt} \rightleftharpoons \text{RasGTP_Golgi_GM}$	
6	ca_bind_rasGRP	ca bind rasGRP	$\text{Ca} + \text{RasGRP_cyt} \rightleftharpoons \text{CaRasGRP1_cyt}$	
7	RasGRP_DAG_GEF	RasGRP_DAG GEF	$\text{RasGDP_Golgi_GM} \xrightleftharpoons{\text{RasGRP_DAG_GM}} \text{RasGTP_Golgi_GM}$	
8	rasGTP_pal- _translocation	rasGTP pal translocation	$\text{RasGTP_pal_cyt} \rightleftharpoons \text{RasGTP_PM}$	
9	PLCg_dephos	PLCg dephos	$\text{PLC_act_PM} \rightleftharpoons \text{PLC_PM}$	
10	basal_GAP	basal GAP	$\text{RasGTP_PM} \rightleftharpoons \text{RasGDP_PM}$	
11	CAPRI- _translocation	CAPRI translocation	$\text{CaCAPRI_cyt} \rightleftharpoons \text{CaCAPRI_PM_PM}$	
12	reaction5	reaction5	$\text{DAG_GM_GM} + \text{RasGRP_cyt} \rightleftharpoons \text{RasGRP_DAG_GM}$	
13	RasGDP_pal	RasGDP pal	$\text{RasGDP_Golgi_GM} \rightleftharpoons \text{RasGDP_pal_cyt}$	
14	CAPRI_GAP	CAPRI GAP	$\text{RasGTP_PM} \xrightleftharpoons{\text{CaCAPRI_PM_PM}} \text{RasGDP_PM}$	
15	RasGDPpal- _translocation	RasGDPpal translocation	$\text{RasGDP_pal_cyt} \rightleftharpoons \text{RasGDP_PM}$	
16	sos_grb2- _binding	sos grb2 binding	$\text{Sos_cyt} + \text{Grb2_cyt} \rightleftharpoons \text{SosGrb2_cyt}$	

Nº	Id	Name	Reaction Equation	SBO
17	RasGDP_depall-translocate	RasGDP depal translocate	$\text{RasGDP_depall_cyt} \rightleftharpoons \text{RasGDP_Golgi_GM}$	
18	Ras_GTP_palm1	Ras GTP palm1	$\text{RasGTP_Golgi_GM} \rightleftharpoons \text{RasGTP_pal_cyt}$	
19	RasPal_basal-GAP	RasPal basal GAP	$\text{RasGTP_pal_cyt} \rightleftharpoons \text{RasGDP_pal_cyt}$	
20	basal_cyt-depal_GEF	basal cyt depal GEF	$\text{RasGTP_depall_cyt} \rightleftharpoons \text{RasGDP_depall_cyt}$	
21	caPLCe_gen_DAG	caPLCe gen DAG	$\text{PIP2_GM_GM} \xrightleftharpoons{\text{Ras_CaPLCe_GM}} \text{DAG_GM_GM} + \text{IP3}$	
22	CaRasGRP_act-RasGM	CaRasGRP act RasGM	$\text{RasGDP_Golgi_GM} \xrightleftharpoons{\text{Ca_RasGRP_GM_GM}} \text{RasGTP_Golgi_GM}$	
23	PIP2_hydrolysis	PIP2 hydrolysis	$\text{PIP2_PM} \xrightleftharpoons{\text{PLC_act_PM}} \text{DAG_PM} + \text{IP3}$	
24	Sos_activation	Sos activation	$\text{SosGrb2_cyt} + \text{Shc_star_PM} \rightleftharpoons \text{SGS_PM}$	
25	PIP_synthesis	PIP synthesis	$\text{PI_PM} \rightleftharpoons \text{PIP_PM}$	
26	EGF-internalization	EGF internalization	$\text{Activated_EGFR_PM} \rightleftharpoons \emptyset$	
27	calcium_buffer	calcium buffer	$\text{buffer_cyt} + \text{Ca} \rightleftharpoons \text{ca_buffer_cyt}$	
28	RasGM_basal_GAP	RasGM basal GAP	$\text{RasGTP_Golgi_GM} \rightleftharpoons \text{RasGDP_Golgi_GM}$	
29	reaction0	reaction0	$\text{Shc_star_PM} \rightleftharpoons \text{Shc_PM}$	
30	EGFR_binding	EGFR binding	$\text{EGF_EC} + \text{EGFR_PM} \rightleftharpoons \text{Activated_EGFR_PM}$	
31	ca_act_PLCe	ca act PLCe	$\text{Ca} + \text{PLCe_cyt} \rightleftharpoons \text{Ca_PLCe_cyt}$	
32	ras_act_PLCe	ras_act.PLCe	$\text{RasGTP_Golgi_GM} + \text{Ca_PLCe_cyt} \rightleftharpoons \text{Ras_CaPLCe_GM}$	
33	RasGDP_depall2	RasGDP depall2	$\text{RasGDP_PM} \rightleftharpoons \text{RasGDP_depall_cyt}$	
34	CaRasGRP-translocation	CaRasGRP translocation	$\text{CaRasGRP1_cyt} \rightleftharpoons \text{Ca_RasGRP_GM_GM}$	
35	reaction2	reaction2	$\text{Rinh} + \text{Ca} \rightleftharpoons \text{RinhCa}$	

Nº	Id	Name	Reaction Equation	SBO
36	EGF_act- _PLCgamma	EGF act PLCgamma	$\text{PLC_PM} \xrightleftharpoons{\text{Activated_EGFR_PM}} \text{PLC_act_PM}$	
37	Ca_binds_IP3R	Ca_binds_IP3R	$\text{Ract} + \text{Ca} \rightleftharpoons \text{RactCa}$	
38	reaction7	reaction7	$\text{DAG_GM_GM} \rightleftharpoons \emptyset$	
39	Sos_act_RasPM	Sos act RasPM	$\text{RasGDP_PM} \xrightleftharpoons{\text{SGS_PM}} \text{RasGTP_PM}$	
40	flux1	flux1	$\text{Ca} \xrightleftharpoons{\text{ER_erMembrane, serca}} \text{Ca_ER}$	
41	flux0	flux0	$\text{Ca} \xrightleftharpoons{\text{ER_erMembrane, RactCa, Ract, IP3, Rinh, RinhCa}} \text{Ca_ER}$	
42	flux2	flux2	$\text{Ca} \xrightleftharpoons{\text{ER_erMembrane}} \text{Ca_ER}$	
43	Ras_PM_depal1	Ras PM depal1	$\text{RasGTP_PM} \rightleftharpoons \text{RasGTP_depal_cyt}$	

7.1 Reaction PIP2_synthesis

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

Name PIP2 synthesis

Reaction equation



Reactant

Table 6: Properties of each reactant.

Id	Name	SBO
PIP_PM		

Product

Table 7: Properties of each product.

Id	Name	SBO
PIP2_PM		

Kinetic Law

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

$$v_1 = (\text{Rate_PIP2Synbasal_PIP2_synthesis} + \text{Rate_PIP2SynStim_PIP2_synthesis}) \cdot [\text{PIP_PM}] \cdot \text{area}(\text{PM}) \quad (14)$$

Table 8: Properties of each parameter.

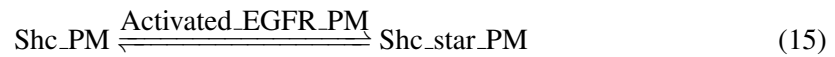
Id	Name	SBO	Value	Unit	Constant
I			0.0	dimensionless · A · m^{-2}	<input checked="" type="checkbox"/>

7.2 Reaction Shc_phosphorylation

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

Name Shc phosphorylation

Reaction equation



Reactant

Table 9: Properties of each reactant.

Id	Name	SBO
Shc_PM		

Modifier

Table 10: Properties of each modifier.

Id	Name	SBO
Activated_EGFR_PM		

Product

Table 11: Properties of each product.

Id	Name	SBO
Shc_star_PM		

Kinetic Law

Derived unit contains undeclared units

$$v_2 = V_{\text{max_Shc_phosphorylation}} \cdot [\text{Shc_PM}] \cdot \frac{1}{K_m + [\text{Shc_PM}]} \cdot \text{area}(\text{PM}) \quad (16)$$

Table 12: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
I			0.0	dimensionless · A · m ⁻²	<input checked="" type="checkbox"/>
Km			1032.0	10 ¹² dimensionless · item · m ⁻²	<input checked="" type="checkbox"/>

7.3 Reaction Ca_bind_CAPRI

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

Name Ca bind CAPRI

Reaction equation



Reactants

Table 13: Properties of each reactant.

Id	Name	SBO
Ca		
CAPRI_cyt		

Product

Table 14: Properties of each product.

Id	Name	SBO
CaCAPRI_cyt		

Kinetic Law

Derived unit contains undeclared units

$$v_3 = (\text{Kf} \cdot 0.00166112956810631 \cdot [\text{Ca}] \cdot 0.00166112956810631 \cdot [\text{CAPRI_cyt}] + ((\text{Kr} \cdot 0.00166112956810631 \cdot [\text{CaCAPRI_cyt}]))) \cdot \text{vol}(\text{cyt}) \cdot 1 \cdot \frac{1}{\text{KMOLE}} \quad (18)$$

Table 15: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
Kf			0.1	1000 dimensionless · m ³ · mol ⁻¹ · s ⁻¹	✓
Kr			0.5	s ⁻¹	✓

7.4 Reaction IP3_degradation

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

Name IP3 degradation

Reaction equation



Reactant

Table 16: Properties of each reactant.

Id	Name	SBO
IP3		

Kinetic Law

Derived unit contains undeclared units

$$v_4 = \text{kIP3deg} \cdot (0.00166112956810631 \cdot [\text{IP3}] + (\text{IP3_basal})) \cdot \text{vol}(\text{cyt}) \cdot 1 \cdot \frac{1}{\text{KMOLE}} \quad (20)$$

Table 17: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
kIP3deg			0.5	s ⁻¹	<input checked="" type="checkbox"/>
IP3_basal			0.0	0.0010 dimensionless · m ⁻³ · mol	<input checked="" type="checkbox"/>

7.5 Reaction RasGTP_depal_translocate

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

Name RasGTP depal translocate

Reaction equation



Reactant

Table 18: Properties of each reactant.

Id	Name	SBO
RasGTP_depal_cyt		

Product

Table 19: Properties of each product.

Id	Name	SBO
RasGTP_Golgi_GM		

Kinetic Law

Derived unit contains undeclared units

$$v_5 = (K_f \cdot 0.00166112956810631 \cdot [\text{RasGTP_depal_cyt}] + ((K_r \cdot [\text{RasGTP_Golgi_GM}]))) \cdot \text{area}(\text{GM}) \quad (22)$$

Table 20: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
I			0.00	dimensionless · A · m ⁻²	✓
Kf			120.00	10 ¹⁵ dimensionless · item · m · mol ⁻¹ · s ⁻¹	✓
Kr			0.01	s ⁻¹	✓

7.6 Reaction `ca_bind_rasGRP`

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

Name `ca bind rasGRP`

Reaction equation



Reactants

Table 21: Properties of each reactant.

Id	Name	SBO
Ca		
RasGRP_cyt		

Product

Table 22: Properties of each product.

Id	Name	SBO
CaRasGRP1_cyt		

Kinetic Law

Derived unit contains undeclared units

$$v_6 = (K_f \cdot 0.00166112956810631 \cdot [\text{Ca}] \cdot 0.00166112956810631 \cdot [\text{RasGRP_cyt}] + ((K_r \cdot 0.00166112956810631 \cdot [\text{CaRasGRP1_cyt}]))) \cdot \text{vol}(\text{cyt}) \cdot 1 \cdot \frac{1}{\text{KMOLE}} \quad (24)$$

Table 23: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
Kf			0.1	1000 dimensionless · m ³ · mol ⁻¹ · s ⁻¹	<input checked="" type="checkbox"/>
Kr			0.5	s ⁻¹	<input checked="" type="checkbox"/>

7.7 Reaction RasGRP_DAG_GEF

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

Name RasGRP_DAG GEF

Reaction equation



Reactant

Table 24: Properties of each reactant.

Id	Name	SBO
RasGDP_Golgi_GM		

Modifier

Table 25: Properties of each modifier.

Id	Name	SBO
RasGRP_DAG_GM		

Product

Table 26: Properties of each product.

Id	Name	SBO
RasGTP_Golgi_GM		

Kinetic Law

Derived unit contains undeclared units

$$\begin{aligned}
 v_7 &= V_{\max_RasGRP_DAG_GEF} \cdot [RasGDP_Golgi_GM] \\
 &\quad \cdot \frac{1}{K_m + [RasGDP_Golgi_GM]} \cdot \text{area (GM)}
 \end{aligned}
 \tag{26}$$

Table 27: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
I			0.0	dimensionless · A · m ⁻²	✓
Km			600.0	10 ¹² dimensionless · item · m ⁻²	✓

7.8 Reaction `rasGTP_pal_translocation`

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

Name rasGTP pal translocation

Reaction equation



Reactant

Table 28: Properties of each reactant.

Id	Name	SBO
RasGTP_pal_cyt		

Product

Table 29: Properties of each product.

Id	Name	SBO
RasGTP_PM		

Kinetic Law

Derived unit contains undeclared units

$$v_8 = (\text{Kf} \cdot 0.00166112956810631 \cdot [\text{RasGTP_pal_cyt}] + ((\text{Kr} \cdot [\text{RasGTP_PM}]))) \cdot \text{area}(\text{PM}) \quad (28)$$

Table 30: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
I			0.00	dimensionless · A · m ⁻²	✓
Kf			120.00	10 ¹⁵ dimensionless · item · m · mol ⁻¹ · s ⁻¹	✓
Kr			0.01	s ⁻¹	✓

7.9 Reaction PLCg_dephos

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

Name PLCg dephos

Reaction equation



Reactant

Table 31: Properties of each reactant.

Id	Name	SBO
PLC_act_PM		

Product

Table 32: Properties of each product.

Id	Name	SBO
PLC_PM		

Kinetic Law

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

$$v_9 = (\text{Kf} \cdot [\text{PLC_act_PM}] + ((\text{Kr} \cdot [\text{PLC_PM}]))) \cdot \text{area}(\text{PM}) \quad (30)$$

Table 33: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
I			0.0	dimensionless · A · m ⁻²	✓
Kf			0.2	s ⁻¹	✓
Kr			0.0	s ⁻¹	✓

7.10 Reaction basal_GAP

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

Name basal GAP

Reaction equation



Reactant

Table 34: Properties of each reactant.

Id	Name	SBO
RasGTP_PM		

Product

Table 35: Properties of each product.

Id	Name	SBO
RasGDP_PM		

Kinetic Law

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

$$v_{10} = (\text{Kf} \cdot [\text{RasGTP_PM}] + ((\text{Kr} \cdot [\text{RasGDP_PM}]))) \cdot \text{area}(\text{PM}) \quad (32)$$

Table 36: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
I			0.000	dimensionless · A · m ⁻²	✓
Kf			10 ⁻⁴	s ⁻¹	✓
Kr			0.000	s ⁻¹	✓

7.11 Reaction CAPRI_translocation

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

Name CAPRI translocation

Reaction equation



Reactant

Table 37: Properties of each reactant.

Id	Name	SBO
CaCAPRI_cyt		

Product

Table 38: Properties of each product.

Id	Name	SBO
CaCAPRI_PM_PM		

Kinetic Law

Derived unit contains undeclared units

$$v_{11} = (K_f \cdot 0.00166112956810631 \cdot [\text{CaCAPRI_cyt}] + ((K_r \cdot [\text{CaCAPRI_PM_PM}]))) \cdot \text{area}(\text{PM}) \quad (34)$$

Table 39: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
I			0.0	dimensionless · A · m ⁻²	✓
Kf			120.0	10 ¹⁵ dimensionless · item · m · mol ⁻¹ · s ⁻¹	✓
Kr			0.1	s ⁻¹	✓

7.12 Reaction `reaction5`

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

Name `reaction5`

Reaction equation



Reactants

Table 40: Properties of each reactant.

Id	Name	SBO
DAG_GM_GM		
RasGRP_cyt		

Product

Table 41: Properties of each product.

Id	Name	SBO
RasGRP_DAG_GM		

Kinetic Law

Derived unit contains undeclared units

$$v_{12} = (K_f \cdot [\text{DAG_GM_GM}] \cdot 0.00166112956810631 \cdot [\text{RasGRP_cyt}] + ((K_r \cdot [\text{RasGRP_DAG_GM}]))) \cdot \text{area (GM)} \quad (36)$$

Table 42: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
I			0.0	dimensionless · A · m ⁻²	✓
Kf			0.5	1000 dimensionless · m ³ · mol ⁻¹ · s ⁻¹	✓
Kr			0.1	s ⁻¹	✓

7.13 Reaction RasGDP_pal

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

Name RasGDP pal

Reaction equation



Reactant

Table 43: Properties of each reactant.

Id	Name	SBO
	RasGDP_Golgi_GM	

Product

Table 44: Properties of each product.

Id	Name	SBO
	RasGDP_pal_cyt	

Kinetic Law

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

$$v_{13} = (\text{Kf} \cdot [\text{RasGDP_Golgi_GM}] + ((\text{Kr} \cdot 0.00166112956810631 \cdot [\text{RasGDP_pal_cyt}]))) \cdot \text{area}(\text{GM}) \quad (38)$$

Table 45: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
I			0.000	dimensionless · A · m ⁻²	<input checked="" type="checkbox"/>
Kf			0.015	s ⁻¹	<input checked="" type="checkbox"/>
Kr			10 ⁻⁵	10 ¹⁵ dimensionless · item · m · mol ⁻¹ · s ⁻¹	<input checked="" type="checkbox"/>

7.14 Reaction CAPRI_GAP

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

Name CAPRI GAP

Reaction equation



Reactant

Table 46: Properties of each reactant.

Id	Name	SBO
RasGTP_PM		

Modifier

Table 47: Properties of each modifier.

Id	Name	SBO
CaCAPRI_PM_PM		

Product

Table 48: Properties of each product.

Id	Name	SBO
RasGDP_PM		

Kinetic Law

Derived unit contains undeclared units

$$v_{14} = V_{\max_CAPRI_GAP} \cdot [RasGTP_PM] \cdot \frac{1}{K_m + [RasGTP_PM]} \cdot \text{area}(\text{PM}) \quad (40)$$

Table 49: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
I			0.0	dimensionless · A · m ⁻²	<input checked="" type="checkbox"/>
K _m			1200.0	10 ¹² dimensionless · item · m ⁻²	<input checked="" type="checkbox"/>

7.15 Reaction RasGDPpal_translocation

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

Name RasGDPpal translocation

Reaction equation



Reactant

Table 50: Properties of each reactant.

Id	Name	SBO
RasGDP_pal_cyt		

Product

Table 51: Properties of each product.

Id	Name	SBO
RasGDP_PM		

Kinetic Law

Derived unit contains undeclared units

$$v_{15} = (\text{Kf} \cdot 0.00166112956810631 \cdot [\text{RasGDP_pal_cyt}] + ((\text{Kr} \cdot [\text{RasGDP_PM}]))) \cdot \text{area}(\text{PM}) \quad (42)$$

Table 52: Properties of each parameter.

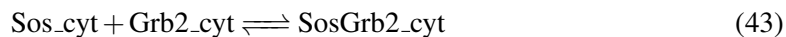
Id	Name	SBO	Value	Unit	Constant
I			0.00	dimensionless · A · m ⁻²	✓
Kf			120.00	10 ¹⁵ dimensionless · item · m · mol ⁻¹ · s ⁻¹	✓
Kr			0.01	s ⁻¹	✓

7.16 Reaction `sos_grb2_binding`

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

Name `sos grb2 binding`

Reaction equation



Reactants

Table 53: Properties of each reactant.

Id	Name	SBO
Sos_cyt		
Grb2_cyt		

Product

Table 54: Properties of each product.

Id	Name	SBO
SosGrb2_cyt		

Kinetic Law

Derived unit contains undeclared units

$$v_{16} = (\text{Kf} \cdot 0.00166112956810631 \cdot [\text{Sos_cyt}] \cdot 0.00166112956810631 \cdot [\text{Grb2_cyt}] + ((\text{Kr} \cdot 0.00166112956810631 \cdot [\text{SosGrb2_cyt}]))) \cdot \text{vol}(\text{cyt}) \cdot 1 \cdot \frac{1}{\text{KMOLE}} \quad (44)$$

Table 55: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
Kf			0.025	1000 dimensionless · m ³ · mol ⁻¹ · s ⁻¹	✓
Kr			0.017	s ⁻¹	✓

7.17 Reaction RasGDP_depal_translocate

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

Name RasGDP depal translocate

Reaction equation



Reactant

Table 56: Properties of each reactant.

Id	Name	SBO
RasGDP_depal_cyt		

Product

Table 57: Properties of each product.

Id	Name	SBO
RasGDP_Golgi_GM		

Kinetic Law

Derived unit contains undeclared units

$$v_{17} = (\text{Kf} \cdot 0.00166112956810631 \cdot [\text{RasGDP_depal_cyt}] + ((\text{Kr} \cdot [\text{RasGDP_Golgi_GM}]))) \cdot \text{area (GM)} \quad (46)$$

Table 58: Properties of each parameter.

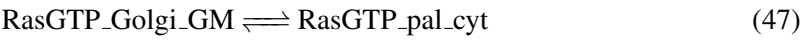
Id	Name	SBO	Value	Unit	Constant
I			0.00	dimensionless · A · m ⁻²	<input checked="" type="checkbox"/>
Kf			120.00	10 ¹⁵ dimensionless · item · m · mol ⁻¹ · s ⁻¹	<input checked="" type="checkbox"/>
Kr			0.01	s ⁻¹	<input checked="" type="checkbox"/>

7.18 Reaction Ras_GTP_palm1

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

Name Ras GTP palm1

Reaction equation



Reactant

Table 59: Properties of each reactant.

Id	Name	SBO
RasGTP_Golgi_GM		

Product

Table 60: Properties of each product.

Id	Name	SBO
RasGTP_pal_cyt		

Kinetic Law

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

$$v_{18} = (Kf \cdot [\text{RasGTP_Golgi_GM}] + ((Kr \cdot 0.00166112956810631 \cdot [\text{RasGTP_pal_cyt}]))) \cdot \text{area}(\text{GAP})$$

(48)

Table 61: Properties of each parameter.

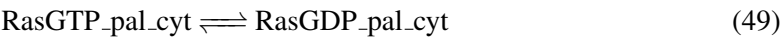
Id	Name	SBO	Value	Unit	Constant
I			0.000	dimensionless · A · m ⁻²	<input checked="" type="checkbox"/>
Kf			$1.58489319246111 \cdot 10^{-4}$	s ⁻¹	<input checked="" type="checkbox"/>
Kr			10^{-5}	10^{15} dimensionless · item · m · mol ⁻¹ · s ⁻¹	<input checked="" type="checkbox"/>

7.19 Reaction RasPal_basal_GAP

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

Name RasPal basal GAP

Reaction equation



Reactant

Table 62: Properties of each reactant.

Id	Name	SBO
RasGTP_pal_cyt		

Product

Table 63: Properties of each product.

Id	Name	SBO
RasGDP_pal_cyt		

Kinetic Law

Derived unit contains undeclared units

$$v_{19} = (K_f \cdot 0.00166112956810631 \cdot [\text{RasGTP_pal_cyt}] + ((K_r \cdot 0.00166112956810631 \cdot [\text{RasGDP_pal_cyt}]))) \cdot \text{vol}(\text{cyt}) \cdot 1 \cdot \frac{1}{\text{KMOLE}} \quad (50)$$

Table 64: Properties of each parameter.

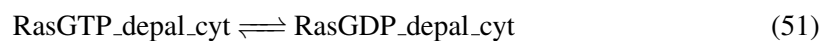
Id	Name	SBO	Value	Unit	Constant
Kf			10^{-4}	s^{-1}	<input checked="" type="checkbox"/>
Kr			0.000	s^{-1}	<input checked="" type="checkbox"/>

7.20 Reaction basal_cyt_depal_GEF

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

Name basal cyt depal GEF

Reaction equation



Reactant

Table 65: Properties of each reactant.

Id	Name	SBO
RasGTP_depall_cyt		

Product

Table 66: Properties of each product.

Id	Name	SBO
RasGDP_depall_cyt		

Kinetic Law

Derived unit contains undeclared units

$$v_{20} = (K_f \cdot 0.00166112956810631 \cdot [\text{RasGTP_depall_cyt}] + ((K_r \cdot 0.00166112956810631 \cdot [\text{RasGDP_depall_cyt}]))) \cdot \text{vol}(\text{cyt}) \cdot 1 \cdot \frac{1}{\text{KMOLE}} \quad (52)$$

Table 67: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
Kf			10 ⁻⁴	s ⁻¹	<input checked="" type="checkbox"/>
Kr			0.000	s ⁻¹	<input checked="" type="checkbox"/>

7.21 Reaction caPLCe_gen_DAG

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

Name caPLCe gen DAG

Reaction equation



Reactant

Table 68: Properties of each reactant.

Id	Name	SBO
PIP2_GM_GM		

Modifier

Table 69: Properties of each modifier.

Id	Name	SBO
Ras_CaPLCe_GM		

Products

Table 70: Properties of each product.

Id	Name	SBO
DAG_GM_GM		
IP3		

Kinetic Law

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

$$v_{21} = k_{\text{act}} \cdot [\text{PIP2_GM_GM}] \cdot [\text{Ras_CaPLCe_GM}] \cdot \text{area}(\text{GM}) \quad (54)$$

Table 71: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
I			0.00	dimensionless · A · m ⁻²	✓
kact			1.18	10 ⁻¹² dimensionless · item ⁻¹ · m ² · s ⁻¹	✓

7.22 Reaction CaRasGRP_act_RasGM

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

Name CaRasGRP act RasGM

Reaction equation



Reactant

Table 72: Properties of each reactant.

Id	Name	SBO
RasGDP_Golgi_GM		

Modifier

Table 73: Properties of each modifier.

Id	Name	SBO
Ca_RasGRP_GM_GM		

Product

Table 74: Properties of each product.

Id	Name	SBO
RasGTP_Golgi_GM		

Kinetic Law

Derived unit contains undeclared units

$$v_{22} = \text{Vmax_CaRasGRP_act_RasGM} \cdot [\text{RasGDP_Golgi_GM}] \cdot \frac{1}{\text{Km} + [\text{RasGDP_Golgi_GM}]} \cdot \text{area(GM)} \quad (56)$$

Table 75: Properties of each parameter.

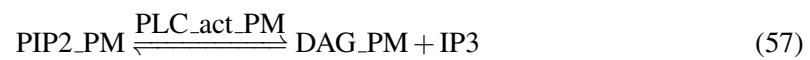
Id	Name	SBO	Value	Unit	Constant
I			0.0	dimensionless · A · m ⁻²	✓
Km			1200.0	10 ¹² dimensionless · item · m ⁻²	✓

7.23 Reaction PIP2_hydrolysis

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

Name PIP2 hydrolysis

Reaction equation



Reactant

Table 76: Properties of each reactant.

Id	Name	SBO
PIP2_PM		

Modifier

Table 77: Properties of each modifier.

Id	Name	SBO
PLC_act_PM		

Products

Table 78: Properties of each product.

Id	Name	SBO
DAG_PM		
IP3		

Kinetic Law

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

$$v_{23} = k_{\text{PIP2hyd}} \cdot [\text{PIP2_PM}] \cdot [\text{PLC_act_PM}] \cdot \text{area}(\text{PM}) \quad (58)$$

Table 79: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
I			0.000	dimensionless · A · m ⁻²	<input checked="" type="checkbox"/>
k_PIP2hyd			1.188	10 ⁻¹² dimensionless · item ⁻¹ · m ² · s ⁻¹	<input checked="" type="checkbox"/>

7.24 Reaction Sos_activation

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

Name Sos activation

Reaction equation



Reactants

Table 80: Properties of each reactant.

Id	Name	SBO
	SosGrb2_cyt	
	Shc_star_PM	

Product

Table 81: Properties of each product.

Id	Name	SBO
	SGS_PM	

Kinetic Law

Derived unit contains undeclared units

$$v_{24} = (\text{Kf} \cdot 0.00166112956810631 \cdot [\text{SosGrb2_cyt}] \cdot [\text{Shc_star_PM}] + ((\text{Kr} \cdot [\text{SGS_PM}]))) \cdot \text{area (PM)} \quad (60)$$

Table 82: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
I			0.0	dimensionless · A · m ⁻²	<input checked="" type="checkbox"/>
Kf			90.0	1000 dimensionless · m ³ · mol ⁻¹ · s ⁻¹	<input checked="" type="checkbox"/>
Kr			0.1	s ⁻¹	<input checked="" type="checkbox"/>

7.25 Reaction PIP_synthesis

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

Name PIP synthesis

Reaction equation



Reactant

Table 83: Properties of each reactant.

Id	Name	SBO
PI_PM		

Product

Table 84: Properties of each product.

Id	Name	SBO
PIP_PM		

Kinetic Law

Derived unit s⁻¹ · item

$$v_{25} = (\text{Ratebasal_PIPsyn_PIP_synthesis} + \text{Rateestim_PIPsyn_PIP_synthesis}) \cdot [\text{PI_PM}] \cdot \text{area}(\text{PM}) \quad (62)$$

Table 85: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
I			0.0	dimensionless · A · m ⁻²	<input checked="" type="checkbox"/>

7.26 Reaction `EGF_internalization`

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

Name `EGF_internalization`

Reaction equation



Reactant

Table 86: Properties of each reactant.

Id	Name	SBO
	Activated_EGFR_PM	

Kinetic Law

Derived unit s⁻¹ · item

$$v_{26} = K_f \cdot [\text{Activated_EGFR_PM}] \cdot \text{area}(\text{PM}) \quad (64)$$

Table 87: Properties of each parameter.

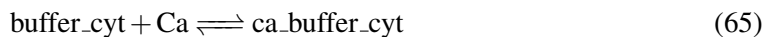
Id	Name	SBO	Value	Unit	Constant
I			0.000	dimensionless · A · m ⁻²	<input checked="" type="checkbox"/>
Kf			0.002	s ⁻¹	<input checked="" type="checkbox"/>
Kr			0.000	10 ¹² dimensionless · item · m ⁻² · s ⁻¹	<input checked="" type="checkbox"/>

7.27 Reaction `calcium_buffer`

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

Name calcium buffer

Reaction equation



Reactants

Table 88: Properties of each reactant.

Id	Name	SBO
	buffer_cyt	
	Ca	

Product

Table 89: Properties of each product.

Id	Name	SBO
	ca_buffer_cyt	

Kinetic Law

Derived unit contains undeclared units

$$v_{27} = (\text{Kf} \cdot 0.00166112956810631 \cdot [\text{buffer_cyt}] \cdot 0.00166112956810631 \cdot [\text{Ca}] + ((\text{Kr} \cdot 0.00166112956810631 \cdot [\text{ca_buffer_cyt}]))) \cdot \text{vol}(\text{cyt}) \cdot 1 \cdot \frac{1}{\text{KMOLE}} \quad (66)$$

Table 90: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
Kf			50.0	1000 dimensionless · m ³ · mol ⁻¹ · s ⁻¹	✓
Kr			10.0	s ⁻¹	✓

7.28 Reaction RasGM_basal_GAP

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

Name RasGM basal GAP

Reaction equation



Reactant

Table 91: Properties of each reactant.

Id	Name	SBO
	RasGTP_Golgi_GM	

Product

Table 92: Properties of each product.

Id	Name	SBO
	RasGDP_Golgi_GM	

Kinetic Law

Derived unit contains undeclared units

$$v_{28} = V_{\max} \cdot [\text{RasGTP_Golgi_GM}] \cdot \frac{1}{K_m + [\text{RasGTP_Golgi_GM}]} \cdot \text{area}(\text{GM}) \quad (68)$$

Table 93: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
I			0.0	dimensionless · A · m ⁻²	✓
Km			600.0	10 ¹² dimensionless · item · m ⁻²	✓
Vmax			1.0	10 ¹² dimensionless · item · m ⁻² · s ⁻¹	✓

7.29 Reaction `reaction0`

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

Name `reaction0`

Reaction equation



Reactant

Table 94: Properties of each reactant.

Id	Name	SBO
Shc_star_PM		

Product

Table 95: Properties of each product.

Id	Name	SBO
Shc_PM		

Kinetic Law

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

$$v_{29} = (\text{Kf} \cdot [\text{Shc_star_PM}] + ((\text{Kr} \cdot [\text{Shc_PM}]))) \cdot \text{area}(\text{PM}) \quad (70)$$

Table 96: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
I			0.0	dimensionless · A · m ⁻²	<input checked="" type="checkbox"/>
Kf			0.5	s ⁻¹	<input checked="" type="checkbox"/>
Kr			0.0	s ⁻¹	<input checked="" type="checkbox"/>

7.30 Reaction EGFR_binding

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

Name EGFR_binding

Reaction equation



Reactants

Table 97: Properties of each reactant.

Id	Name	SBO
EGF_EC		
EGFR_PM		

Product

Table 98: Properties of each product.

Id	Name	SBO
Activated_EGFR_PM		

Kinetic Law

Derived unit contains undeclared units

$$v_{30} = (K_f \cdot 0.00166112956810631 \cdot [\text{EGF_EC}] \cdot [\text{EGFR_PM}] + ((K_r \cdot [\text{Activated_EGFR_PM}]))) \cdot \text{area}(\text{PM}) \quad (72)$$

Table 99: Properties of each parameter.

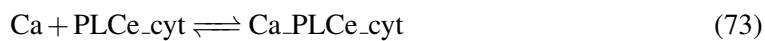
Id	Name	SBO	Value	Unit	Constant
I			0.00	dimensionless · A · m ⁻²	✓
Kf			1.00	1000 dimensionless · m ³ · mol ⁻¹ · s ⁻¹	✓
Kr			0.01	s ⁻¹	✓

7.31 Reaction `ca_act_PLCe`

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

Name `ca act PLCe`

Reaction equation



Reactants

Table 100: Properties of each reactant.

Id	Name	SBO
Ca		
PLCe_cyt		

Product

Table 101: Properties of each product.

Id	Name	SBO
Ca_PLCe_cyt		

Kinetic Law

Derived unit contains undeclared units

$$v_{31} = (K_f \cdot 0.00166112956810631 \cdot [\text{Ca}] \cdot 0.00166112956810631 \cdot [\text{PLCe_cyt}] + ((K_r \cdot 0.00166112956810631 \cdot [\text{Ca_PLCe_cyt}]))) \cdot \text{vol}(\text{cyt}) \cdot 1 \cdot \frac{1}{\text{KMOLE}} \quad (74)$$

Table 102: Properties of each parameter.

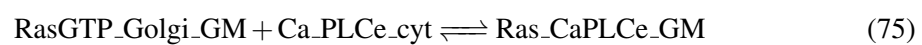
Id	Name	SBO	Value	Unit	Constant
Kf			3.0	1000 dimensionless · m ³ · mol ⁻¹ · s ⁻¹	<input checked="" type="checkbox"/>
Kr			1.0	s ⁻¹	<input checked="" type="checkbox"/>

7.32 Reaction `ras_act_PLCe`

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

Name `ras_act_PLCe`

Reaction equation



Reactants

Table 103: Properties of each reactant.

Id	Name	SBO
RasGTP_Golgi_GM		
Ca_PLCE_cyt		

Product

Table 104: Properties of each product.

Id	Name	SBO
Ras_CaPLCE_GM		

Kinetic Law

Derived unit contains undeclared units

$$v_{32} = (K_f \cdot [\text{RasGTP_Golgi_GM}] \cdot 0.00166112956810631 \cdot [\text{Ca_PLCE_cyt}] + ((K_r \cdot [\text{Ras_CaPLCE_GM}]))) \cdot \text{area (GM)} \quad (76)$$

Table 105: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
I			0.0	dimensionless · A · m ⁻²	✓
Kf			15.0	1000 dimensionless · m ³ · mol ⁻¹ · s ⁻¹	✓
Kr			1.0	s ⁻¹	✓

7.33 Reaction RasGDP_depal2

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

Name RasGDP depal2

Reaction equation



Reactant

Table 106: Properties of each reactant.

Id	Name	SBO
RasGDP_PM		

Product

Table 107: Properties of each product.

Id	Name	SBO
RasGDP_depal_cyt		

Kinetic Law

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

$$v_{33} = (\text{Kf} \cdot [\text{RasGDP_PM}] + ((\text{Kr} \cdot 0.00166112956810631 \cdot [\text{RasGDP_depal_cyt}]))) \cdot \text{area}(\text{PM}) \quad (78)$$

Table 108: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
I			0.000	dimensionless · A · m ⁻²	✓
Kf			10 ⁻⁴	s ⁻¹	✓
Kr			0.000	10 ¹⁵ dimensionless · item · m · mol ⁻¹ · s ⁻¹	✓

7.34 Reaction CaRasGRP_translocation

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

Name CaRasGRP translocation

Reaction equation



Reactant

Table 109: Properties of each reactant.

Id	Name	SBO
CaRasGRP1_cyt		

Product

Table 110: Properties of each product.

Id	Name	SBO
Ca_RasGRP_GM_GM		

Kinetic Law

Derived unit contains undeclared units

$$v_{34} = (K_f \cdot 0.00166112956810631 \cdot [\text{CaRasGRP1_cyt}] + ((K_r \cdot [\text{Ca_RasGRP_GM_GM}]))) \cdot \text{area (GM)} \quad (80)$$

Table 111: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
I			0.0	dimensionless · A · m ⁻²	✓
Kf			10.0	10 ¹⁵ dimensionless · item · m · mol ⁻¹ · s ⁻¹	✓
Kr			5.0	s ⁻¹	✓

7.35 Reaction `reaction2`

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

Name `reaction2`

Reaction equation



Reactants

Table 112: Properties of each reactant.

Id	Name	SBO
Rinh		
Ca		

Product

Table 113: Properties of each product.

Id	Name	SBO
RinhCa		

Kinetic Law

Derived unit contains undeclared units

$$v_{35} = (\text{Kon_reaction2} \cdot [\text{Rinh}] \cdot 0.00166112956810631 \cdot [\text{Ca}] + ((\text{Koff_reaction2} \cdot [\text{RinhCa}]))) \cdot \text{area}(\text{erMembrane}) \quad (82)$$

Table 114: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
I			0.0	dimensionless · A · m ⁻²	<input checked="" type="checkbox"/>

7.36 Reaction [EGF_act_PLCgamma](#)

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

Name EGF act PLCgamma

Reaction equation



Reactant

Table 115: Properties of each reactant.

Id	Name	SBO
PLC_PM		

Modifier

Table 116: Properties of each modifier.

Id	Name	SBO
Activated_EGFR_PM		

Product

Table 117: Properties of each product.

Id	Name	SBO
PLC_act_PM		

Kinetic Law

Derived unit contains undeclared units

$$v_{36} = V_{\max_EGF_act_PLC\gamma} \cdot [PLC_PM] \cdot \frac{1}{K_m + [PLC_PM]} \cdot \text{area}(PM) \quad (84)$$

Table 118: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
I			0.0	dimensionless · A · m ⁻²	✓
Km			410.0	10 ¹² dimensionless · item · m ⁻²	✓

7.37 Reaction Ca_binds_IP3R

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

Name Ca_binds_IP3R

Reaction equation



Reactants

Table 119: Properties of each reactant.

Id	Name	SBO
Ract		
Ca		

Product

Table 120: Properties of each product.

Id	Name	SBO
RactCa		

Kinetic Law

Derived unit contains undeclared units

$$v_{37} = ([\text{Ract}] \cdot K_f \cdot 0.00166112956810631 \cdot [\text{Ca}] + ((K_r.\text{Ca_binds_IP3R} \cdot [\text{RactCa}]))) \cdot \text{area}(\text{erMembrane}) \quad (86)$$

Table 121: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
I			0.0	dimensionless · A · m ⁻²	✓
Kf			1000.0	1000 dimensionless · m ³ · mol ⁻¹ · s ⁻¹	✓

7.38 Reaction `reaction7`

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

Name `reaction7`

Reaction equation



Reactant

Table 122: Properties of each reactant.

Id	Name	SBO
DAG_GM_GM		

Kinetic Law

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

$$v_{38} = K_f \cdot [\text{DAG_GM_GM}] \cdot \text{area}(\text{GM}) \quad (88)$$

Table 123: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
I			0.00	dimensionless · A · m ⁻²	✓
Kf			0.25	s ⁻¹	✓
Kr			0.00	10 ¹² dimensionless · item · m ⁻² · s ⁻¹	✓

7.39 Reaction Sos_act_RasPM

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

Name Sos act RasPM

Reaction equation



Reactant

Table 124: Properties of each reactant.

Id	Name	SBO
RasGDP_PM		

Modifier

Table 125: Properties of each modifier.

Id	Name	SBO
SGS_PM		

Product

Table 126: Properties of each product.

Id	Name	SBO
RasGTP_PM		

Kinetic Law

Derived unit contains undeclared units

$$v_{39} = V_{\max_Sos_act_RasPM} \cdot [RasGDP_PM] \cdot \frac{1}{K_m + [RasGDP_PM]} \cdot \text{area}(PM) \quad (90)$$

Table 127: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
I			0.0	dimensionless · A · m ⁻²	✓
Km			600.0	10 ¹² dimensionless · item · m ⁻²	✓

7.40 Reaction `flux1`

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

Name `flux1`

Reaction equation



Reactant

Table 128: Properties of each reactant.

Id	Name	SBO
Ca		

Modifiers

Table 129: Properties of each modifier.

Id	Name	SBO
ER_erMembrane		
serca		

Product

Table 130: Properties of each product.

Id	Name	SBO
Ca_ER		

Kinetic Law

Derived unit contains undeclared units

$$\begin{aligned}
 v_{40} = & [\text{ER_erMembrane}] \cdot [\text{serca}] \cdot vP \cdot 0.00166112956810631 \cdot [\text{Ca}] \cdot 0.00166112956810631 \\
 & \cdot [\text{Ca}] \cdot \frac{1}{kP \cdot kP + 0.00166112956810631 \cdot [\text{Ca}] \cdot 0.00166112956810631 \cdot [\text{Ca}]} \\
 & \cdot \text{area(erMembrane)} \cdot 1 \cdot \frac{1}{\text{KMOLE}}
 \end{aligned}
 \tag{92}$$

Table 131: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
I			0.000	dimensionless · A · m ⁻²	✓
vP			0.066	10 ⁻³³ dimensionless · item ⁻² · m ² · mol · s ⁻¹	✓
kP			0.270		✓

7.41 Reaction flux0

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

Name flux0

Reaction equation



Reactant

Table 132: Properties of each reactant.

Id	Name	SBO
Ca		

Modifiers

Table 133: Properties of each modifier.

Id	Name	SBO
ER_erMembrane		
RactCa		
Ract		
IP3		
Rinh		
RinhCa		

Product

Table 134: Properties of each product.

Id	Name	SBO
Ca_ER		

Kinetic Law

Derived unit contains undeclared units

$$\begin{aligned}
v_{41} = & \left(0.25 \cdot [\text{ER_erMembrane}] \cdot ([\text{RactCa}] + [\text{Ract}]) \right. \\
& \cdot (0.00166112956810631 \cdot [\text{Ca_ER}] + ((0.00166112956810631 \cdot [\text{Ca}]))) \\
& \cdot \left(0.00166112956810631 \cdot [\text{IP3}] \cdot [\text{RactCa}] \cdot [\text{Rinh}] \cdot \frac{1}{0.00166112956810631 \cdot [\text{IP3}] + \text{dI}} \right. \\
& \quad \cdot \frac{1}{[\text{RactCa}] + [\text{Ract}]} \cdot \frac{1}{[\text{RinhCa}] + [\text{Rinh}]} \Big)^3 \cdot \text{singleChanFlux} \Big) \\
& \cdot \text{area(erMembrane)} \cdot 1 \cdot \frac{1}{\text{KMOLE}}
\end{aligned}
\tag{94}$$

Table 135: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
I			0.00	dimensionless · A · m ⁻²	✓
dI			0.80	0.0010 dimensionless · m ⁻³ · mol	✓
singleChanFlux			4.69	10 ⁻³⁰ dimensionless · item ⁻² · m ⁵ · s ⁻¹	✓

7.42 Reaction flux2

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

Name flux2

Reaction equation



Reactant

Table 136: Properties of each reactant.

Id	Name	SBO
Ca		

Modifier

Table 137: Properties of each modifier.

Id	Name	SBO
ER_erMembrane		

Product

Table 138: Properties of each product.

Id	Name	SBO
Ca_ER		

Kinetic Law

Derived unit contains undeclared units

$$v_{42} = ([ER_erMembrane] \cdot (0.00166112956810631 \cdot [Ca_ER] + ((0.00166112956810631 \cdot [Ca]))) \cdot vL) \cdot area(erMembrane) \cdot 1 \cdot \frac{1}{KMOLE} \quad (96)$$

Table 139: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
I			0.000	dimensionless · A · m ⁻²	<input checked="" type="checkbox"/>
vL			3.16 · 10 ⁻⁵	10 ⁻¹⁸ dimensionless · item ⁻¹ · m ³ · s ⁻¹	<input checked="" type="checkbox"/>

7.43 Reaction Ras_PM_depal1

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

Name Ras PM depal1

Reaction equation



Reactant

Table 140: Properties of each reactant.

Id	Name	SBO
RasGTP_PM		

Product

Table 141: Properties of each product.

Id	Name	SBO
RasGTP_depal_cyt		

Kinetic Law

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

$$v_{43} = (\text{Kf} \cdot [\text{RasGTP_PM}] + ((\text{Kr} \cdot 0.00166112956810631 \cdot [\text{RasGTP_depal_cyt}]))) \cdot \text{area}(\text{PM}) \quad (98)$$

Table 142: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
I			0.000	dimensionless · A · m ⁻²	✓
Kf			10 ⁻⁴	s ⁻¹	✓
Kr			0.000	10 ¹⁵ dimensionless · item · m · mol ⁻¹ · s ⁻¹	✓

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 `spatialDimensions` > 0 for certain species.

8.1 Species `RasGTP_Golgi_GM`

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

This species takes part in six reactions (as a reactant in `Ras_GTP_palm1`, `RasGM_basal_GAP`, `ras_act_PLCe` and as a product in `RasGTP_depall_translocate`, `RasGRP_DAG_GEF`, `CaRasGRP_act_RasGM`).

$$\frac{d}{dt}\text{RasGTP_Golgi_GM} = v_5 + v_7 + v_{22} - v_{18} - v_{28} - v_{32} \quad (99)$$

8.2 Species `EGF_EC`

Initial concentration $4.02136 \text{ item} \cdot \mu\text{m}^{-3}$

This species takes part in one reaction (as a reactant in `EGFR_binding`), which does not influence its rate of change because this species is on the boundary of the reaction system:

$$\frac{d}{dt}\text{EGF_EC} = 0 \quad (100)$$

8.3 Species `CAPRI_cyt`

Initial concentration $30.1 \text{ item} \cdot \mu\text{m}^{-3}$

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

$$\frac{d}{dt}\text{CAPRI_cyt} = -v_3 \quad (101)$$

8.4 Species `serca`

Initial concentration $45 \text{ item} \cdot \mu\text{m}^{-2}$

This species takes part in one reaction (as a modifier in `flux1`).

$$\frac{d}{dt}\text{serca} = 0 \quad (102)$$

8.5 Species `PIP_PM`

Initial concentration $2857 \text{ item} \cdot \mu\text{m}^{-2}$

This species takes part in two reactions (as a reactant in `PIP2_synthesis` and as a product in `PIP_synthesis`).

$$\frac{d}{dt}\text{PIP_PM} = v_{25} - v_1 \quad (103)$$

8.6 Species `PI_PM`

Initial concentration 142857 item · μm⁻²

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

$$\frac{d}{dt}PI_PM = -v_{25} \quad (104)$$

8.7 Species `Shc_PM`

Initial concentration 186 item · μm⁻²

This species takes part in two reactions (as a reactant in `Shc_phosphorylation` and as a product in `reaction0`).

$$\frac{d}{dt}Shc_PM = v_{29} - v_2 \quad (105)$$

8.8 Species `CaCAPRI_PM_PM`

Initial concentration 0 item · μm⁻²

This species takes part in two reactions (as a product in `CAPRI_translocation` and as a modifier in `CAPRI_GAP`).

$$\frac{d}{dt}CaCAPRI_PM_PM = v_{11} \quad (106)$$

8.9 Species `RactCa`

Initial concentration 2.264 item · μm⁻²

This species takes part in two reactions (as a product in `Ca_binds_IP3R` and as a modifier in `flux0`).

$$\frac{d}{dt}RactCa = v_{37} \quad (107)$$

8.10 Species `Shc_star_PM`

Initial concentration 0 item · μm⁻²

This species takes part in three reactions (as a reactant in `Sos_activation`, `reaction0` and as a product in `Shc_phosphorylation`).

$$\frac{d}{dt}Shc_star_PM = v_2 - v_{24} - v_{29} \quad (108)$$

8.11 Species [EGFR_PM](#)

Initial concentration 21 item · μm⁻²

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

$$\frac{d}{dt}\text{EGFR_PM} = -v_{30} \quad (109)$$

8.12 Species [PLC_act_PM](#)

Initial concentration 0 item · μm⁻²

This species takes part in three reactions (as a reactant in [PLCg_dephos](#) and as a product in [EGF_act_PLCgamma](#) and as a modifier in [PIP2_hydrolysis](#)).

$$\frac{d}{dt}\text{PLC_act_PM} = v_{36} - v_9 \quad (110)$$

8.13 Species [RasGTP_pal_cyt](#)

Initial concentration 0 item · μm⁻³

This species takes part in three reactions (as a reactant in [rasGTP_pal_translocation](#), [RasPal-basal_GAP](#) and as a product in [Ras_GTP_palm1](#)).

$$\frac{d}{dt}\text{RasGTP_pal_cyt} = v_{18} - v_8 - v_{19} \quad (111)$$

8.14 Species [PLC_PM](#)

Initial concentration 100 item · μm⁻²

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

$$\frac{d}{dt}\text{PLC_PM} = v_9 - v_{36} \quad (112)$$

8.15 Species [PIP2_PM](#)

Initial concentration 1072 item · μm⁻²

This species takes part in two reactions (as a reactant in [PIP2_hydrolysis](#) and as a product in [PIP2_synthesis](#)).

$$\frac{d}{dt}\text{PIP2_PM} = v_1 - v_{23} \quad (113)$$

8.16 Species Activated_EGFR_PM

Initial concentration 0 item · μm⁻²

This species takes part in four reactions (as a reactant in [EGF_internalization](#) and as a product in [EGFR_binding](#) and as a modifier in [Shc_phosphorylation](#), [EGF_act_PLCgamma](#)).

$$\frac{d}{dt}\text{Activated_EGFR_PM} = v_{30} - v_{26} \quad (114)$$

8.17 Species Ca

Initial concentration 0 item · μm⁻³

This species takes part in nine reactions (as a reactant in [Ca_bind_CAPRI](#), [ca_bind_rasGRP](#), [calcium_buffer](#), [ca_act_PLCe](#), [reaction2](#), [Ca_binds_IP3R](#), [flux1](#), [flux0](#), [flux2](#)).

$$\frac{d}{dt}\text{Ca} = -v_3 - v_6 - v_{27} - v_{31} - v_{35} - v_{37} - v_{40} - v_{41} - v_{42} \quad (115)$$

8.18 Species Ract

Initial concentration 9.056 item · μm⁻²

This species takes part in two reactions (as a reactant in [Ca_binds_IP3R](#) and as a modifier in [flux0](#)).

$$\frac{d}{dt}\text{Ract} = -v_{37} \quad (116)$$

8.19 Species Rinh

Initial concentration 7.7825 item · μm⁻²

This species takes part in two reactions (as a reactant in [reaction2](#) and as a modifier in [flux0](#)).

$$\frac{d}{dt}\text{Rinh} = -v_{35} \quad (117)$$

8.20 Species RinhCa

Initial concentration 3.5375 item · μm⁻²

This species takes part in two reactions (as a product in [reaction2](#) and as a modifier in [flux0](#)).

$$\frac{d}{dt}\text{RinhCa} = v_{35} \quad (118)$$

8.21 Species IP3

Initial concentration 0 item · μm⁻³

This species takes part in four reactions (as a reactant in [IP3_degradation](#) and as a product in [caPLCe_gen_DAG](#), [PIP2_hydrolysis](#) and as a modifier in [flux0](#)).

$$\frac{d}{dt} \text{IP3} = v_{21} + v_{23} - v_4 \quad (119)$$

8.22 Species RasGDP_Golgi_GM

Initial concentration 50 item · μm⁻²

This species takes part in five reactions (as a reactant in [RasGRP_DAG_GEF](#), [RasGDP_pal](#), [CaRasGRP_act_RasGM](#) and as a product in [RasGDP_depall_translocate](#), [RasGM_basal_GAP](#)).

$$\frac{d}{dt} \text{RasGDP_Golgi_GM} = v_{17} + v_{28} - v_7 - v_{13} - v_{22} \quad (120)$$

8.23 Species Ca_RasGRP_GM_GM

Initial concentration 0 item · μm⁻²

This species takes part in two reactions (as a product in [CaRasGRP_translocation](#) and as a modifier in [CaRasGRP_act_RasGM](#)).

$$\frac{d}{dt} \text{Ca_RasGRP_GM_GM} = v_{34} \quad (121)$$

8.24 Species DAG_GM_GM

Initial concentration 0 item · μm⁻²

This species takes part in three reactions (as a reactant in [reaction5](#), [reaction7](#) and as a product in [caPLCe_gen_DAG](#)).

$$\frac{d}{dt} \text{DAG_GM_GM} = v_{21} - v_{12} - v_{38} \quad (122)$$

8.25 Species RasGRP_DAG_GM

Initial concentration 0 item · μm⁻²

This species takes part in two reactions (as a product in [reaction5](#) and as a modifier in [RasGRP_DAG_GEF](#)).

$$\frac{d}{dt} \text{RasGRP_DAG_GM} = v_{12} \quad (123)$$

8.26 Species CaCAPRI_cyt

Initial concentration 0 item · μm⁻³

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

$$\frac{d}{dt}\text{CaCAPRI_cyt} = v_3 - v_{11} \quad (124)$$

8.27 Species DAG_PM

Initial concentration 2000 item · μm⁻²

This species takes part in one reaction (as a product in [PIP2_hydrolysis](#)).

$$\frac{d}{dt}\text{DAG_PM} = v_{23} \quad (125)$$

8.28 Species RasGTP_depall_cyt

Initial concentration 0 item · μm⁻³

This species takes part in three reactions (as a reactant in [RasGTP_depall_translocate](#), [basal_cyt_depall_GEF](#) and as a product in [Ras_PM_depall1](#)).

$$\frac{d}{dt}\text{RasGTP_depall_cyt} = v_{43} - v_5 - v_{20} \quad (126)$$

8.29 Species RasGDP_depall_cyt

Initial concentration 0 item · μm⁻³

This species takes part in three reactions (as a reactant in [RasGDP_depall_translocate](#) and as a product in [basal_cyt_depall_GEF](#), [RasGDP_depall2](#)).

$$\frac{d}{dt}\text{RasGDP_depall_cyt} = v_{20} + v_{33} - v_{17} \quad (127)$$

8.30 Species RasGDP_pal_cyt

Initial concentration 0 item · μm⁻³

This species takes part in three reactions (as a reactant in [RasGDPpal_translocation](#) and as a product in [RasGDP_pal](#), [RasPal_basal_GAP](#)).

$$\frac{d}{dt}\text{RasGDP_pal_cyt} = v_{13} + v_{19} - v_{15} \quad (128)$$

8.31 Species `Ca_PLCe_cyt`

Initial concentration 0 item · μm^{-3}

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

$$\frac{d}{dt}\text{Ca_PLCe_cyt} = v_{31} - v_{32} \quad (129)$$

8.32 Species `Ras_CaPLCe_GM`

Initial concentration 0 item · μm^{-2}

This species takes part in two reactions (as a product in `ras_act_PLCe` and as a modifier in `caPLCe_gen_DAG`).

$$\frac{d}{dt}\text{Ras_CaPLCe_GM} = v_{32} \quad (130)$$

8.33 Species `PIP2_GM_GM`

Initial concentration 250 item · μm^{-2}

This species takes part in one reaction (as a reactant in `caPLCe_gen_DAG`), which does not influence its rate of change because this species is on the boundary of the reaction system:

$$\frac{d}{dt}\text{PIP2_GM_GM} = 0 \quad (131)$$

8.34 Species `ER_erMembrane`

Initial concentration 2 item · μm^{-2}

This species takes part in three reactions (as a modifier in `flux1`, `flux0`, `flux2`), which do not influence its rate of change because this species is on the boundary of the reaction system:

$$\frac{d}{dt}\text{ER_erMembrane} = 0 \quad (132)$$

8.35 Species `Ca_ER`

Initial concentration 120400 item · μm^{-3}

This species takes part in three reactions (as a product in `flux1`, `flux0`, `flux2`).

$$\frac{d}{dt}\text{Ca_ER} = v_{40} + v_{41} + v_{42} \quad (133)$$

8.36 Species `Sos_cyt`

Initial concentration 81.27 item · μm⁻³

This species takes part in one reaction (as a reactant in `sos_grb2_binding`).

$$\frac{d}{dt} \text{Sos_cyt} = -v_{16} \quad (134)$$

8.37 Species `Grb2_cyt`

Initial concentration 502.67 item · μm⁻³

This species takes part in one reaction (as a reactant in `sos_grb2_binding`).

$$\frac{d}{dt} \text{Grb2_cyt} = -v_{16} \quad (135)$$

8.38 Species `PLCe_cyt`

Initial concentration 12.04 item · μm⁻³

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

$$\frac{d}{dt} \text{PLCe_cyt} = -v_{31} \quad (136)$$

8.39 Species `buffer_cyt`

Initial concentration 120.4 item · μm⁻³

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

$$\frac{d}{dt} \text{buffer_cyt} = -v_{27} \quad (137)$$

8.40 Species `ca_buffer_cyt`

Initial concentration 0 item · μm⁻³

This species takes part in one reaction (as a product in `calcium_buffer`).

$$\frac{d}{dt} \text{ca_buffer_cyt} = v_{27} \quad (138)$$

8.41 Species `SosGrb2_cyt`

Initial concentration 99.33 item · μm⁻³

This species takes part in two reactions (as a reactant in `Sos_activation` and as a product in `sos_grb2_binding`).

$$\frac{d}{dt} \text{SosGrb2_cyt} = v_{16} - v_{24} \quad (139)$$

8.42 Species `SGS_PM`

Initial concentration 0 item · μm^{-2}

This species takes part in two reactions (as a product in `Sos_activation` and as a modifier in `Sos_act_RasPM`).

$$\frac{d}{dt}\text{SGS_PM} = v_{24} \quad (140)$$

8.43 Species `RasGTP_PM`

Initial concentration 0 item · μm^{-2}

This species takes part in five reactions (as a reactant in `basal_GAP`, `CAPRI_GAP`, `Ras_PM-depal1` and as a product in `rasGTP_pal_translocation`, `Sos_act_RasPM`).

$$\frac{d}{dt}\text{RasGTP_PM} = v_8 + v_{39} - v_{10} - v_{14} - v_{43} \quad (141)$$

8.44 Species `RasGDP_PM`

Initial concentration 400 item · μm^{-2}

This species takes part in five reactions (as a reactant in `RasGDP_depal2`, `Sos_act_RasPM` and as a product in `basal_GAP`, `CAPRI_GAP`, `RasGDPpal_translocation`).

$$\frac{d}{dt}\text{RasGDP_PM} = v_{10} + v_{14} + v_{15} - v_{33} - v_{39} \quad (142)$$

8.45 Species `RasGRP_cyt`

Initial concentration 30.1 item · μm^{-3}

This species takes part in two reactions (as a reactant in `ca_bind_rasGRP`, `reaction5`).

$$\frac{d}{dt}\text{RasGRP_cyt} = -v_6 - v_{12} \quad (143)$$

8.46 Species `CaRasGRP1_cyt`

Initial concentration 0 item · μm^{-3}

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

$$\frac{d}{dt}\text{CaRasGRP1_cyt} = v_6 - v_{34} \quad (144)$$

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