

SBML Model Report

Model name: “Fallon2000 - Interleukin-2 dynamics”



May 17, 2018

1 General Overview

This is a document in SBML Level 2 Version 4 format. This model was created by the following two authors: Matthew Grant Roberts¹ and Catherine Lloyd² at June 25th 2010 at noon. and last time modified at June 25th 2010 at noon. Table 1 gives an overview of the quantities of all components of this model.

Table 1: Number of components in this model, which are described in the following sections.

Element	Quantity	Element	Quantity
compartment types	0	compartments	1
species types	0	species	8
events	0	constraints	0
reactions	13	function definitions	5
global parameters	13	unit definitions	6
rules	4	initial assignments	0

Model Notes

This a model from the article:

Computational model for effects of ligand/receptor binding properties on interleukin-2 trafficking dynamics and T cell proliferation response.

Fallon EM, Lauffenburger DA. *Biotechnol Prog* 2000 Sep-Oct;16(5):905-16 [11027188](#) ,

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Abstract:

Multisubunit cytokine receptors such as the heterotrimeric receptor for interleukin-2 (IL-2) are ubiquitous in hematopoietic cell types of importance in biotechnology and are crucial regulators of cell proliferation and differentiation behavior. Dynamics of cytokine/receptor endocytic trafficking can significantly impact cell responses through effects of receptor down-regulation and ligand depletion, and in turn are governed by ligand/receptor binding properties. We describe here a computational model for trafficking dynamics of the IL-2 receptor (IL-2R) system, which is able to predict T cell proliferation responses to IL-2. This model comprises kinetic equations describing binding, internalization, and postendocytic sorting of IL-2 and IL-2R, including an experimentally derived dependence of cell proliferation rate on these properties. Computational results from this model predict that IL-2 depletion can be reduced by decreasing its binding affinity for the IL-2R beta gamma subunit relative to the alpha subunit at endosomal pH, as a result of enhanced ligand sorting to recycling vis-a-vis degradation, and that an IL-2 analogue with such altered binding properties should exhibit increased potency for stimulating the T cell proliferation response. These results are in agreement with our recent experimental findings for the IL-2 analogue termed 2D1 [Fallon, E. M. et al. J. Biol. Chem. 2000, 275, 6790-6797]. Thus, this type of model may enable prediction of beneficial cytokine/receptor binding properties to aid development of molecular design criteria for improvements in applications such as in vivo cytokine therapies and in vitro hematopoietic cell bioreactors.

This model was taken from the [CellML repository](#) and automatically converted to SBML. The original model was: [Fallon EM, Lauffenburger DA. \(2000\) - version=1.0](#)
The original CellML model was created by:

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To cite BioModels Database, please use: [Li C, Donizelli M, Rodriguez N, Dharuri H, Endler L, Chelliah V, Li L, He E, Henry A, Stefan MI, Snoep JL, Hucka M, Le Novre N, Laibe C \(2010\) BioModels Database: An enhanced, curated and annotated resource for published quantitative kinetic models. BMC Syst Biol., 4:92.](#)

2 Unit Definitions

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

2.1 Unit `time`

Name `time`

Definition `60 s`

2.2 Unit `unit_0`

Name `1/(0.0166667*s)`

Definition $(0.0166667 \text{ s})^{-1}$

2.3 Unit `unit_1`

Name `0.06*nl/(mol*s)`

Definition $0.06 \text{ nl} \cdot \text{mol}^{-1} \cdot \text{s}^{-1}$

2.4 Unit `unit_2`

Name `1`

Definition `1`

2.5 Unit `unit_3`

Name `1/Tmol`

Definition Tmol^{-1}

2.6 Unit `unit_4`

Name `1`

Definition `dimensionless0`

2.7 Unit `substance`

Notes Mole is the predefined SBML unit for substance.

Definition `mol`

2.8 Unit volume

Notes Litre is the predefined SBML unit for volume.

Definition 1

2.9 Unit area

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

Definition m^2

2.10 Unit length

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

Definition m

3 Compartment

This model contains one compartment.

Table 2: Properties of all compartments.

Id	Name	SBO	Spatial Dimensions	Size	Unit	Constant	Outside
COMpartment	compartment		3	1	litre	<input checked="" type="checkbox"/>	

3.1 Compartment COMpartment

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

Name compartment

4 Species

This model contains eight 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 Condi- tion
Rs_0	Rs	COMpartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
Cs_0	Cs	COMpartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
Ri_0	Ri	COMpartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
Ci_0	Ci	COMpartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
Li_0	Li	COMpartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
Ld_0	Ld	COMpartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
L_0	L	COMpartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
Y_0	Y	COMpartment	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>

5 Parameters

This model contains 13 global parameters.

Table 4: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
kr	kr		0.014	$(0.0166667 \text{ s})^{-1}$	<input checked="" type="checkbox"/>
kf	kf		0.001	$0.06 \text{ nl} \cdot \text{mol}^{-1} \cdot \text{s}^{-1}$	<input type="checkbox"/>
kre	kre		0.110	$(0.0166667 \text{ s})^{-1}$	<input type="checkbox"/>
kfe	kfe		$1.104 \cdot 10^{-4}$	$0.06 \text{ nl} \cdot \text{mol}^{-1} \cdot \text{s}^{-1}$	<input type="checkbox"/>
kt	kt		0.007	$(0.0166667 \text{ s})^{-1}$	<input checked="" type="checkbox"/>
Vs	Vs		11.000	$(0.0166667 \text{ s})^{-1}$	<input checked="" type="checkbox"/>
ksyn	ksyn		0.001	$(0.0166667 \text{ s})^{-1}$	<input checked="" type="checkbox"/>
ke	ke		0.040	$(0.0166667 \text{ s})^{-1}$	<input checked="" type="checkbox"/>
kx	kx		0.150	$(0.0166667 \text{ s})^{-1}$	<input checked="" type="checkbox"/>
kh	kh		0.035	$(0.0166667 \text{ s})^{-1}$	<input checked="" type="checkbox"/>
Ve	Ve		10^{-14}	1	<input checked="" type="checkbox"/>
NA	NA		$6.022 \cdot 10^{11}$	Tmol^{-1}	<input checked="" type="checkbox"/>
IL2	IL2		1.000	dimensionless ⁰	<input checked="" type="checkbox"/>

6 Function definitions

This is an overview of five function definitions.

6.1 Function definition `Constant_flux_irreversible`

Name Constant flux (irreversible)

Argument v

Mathematical Expression

$$v \quad (1)$$

6.2 Function definition `modifier_function_1`

Name modifier function 1

Arguments parameter, modifier, substrate

Mathematical Expression

$$\text{parameter} \cdot \text{modifier} \cdot \text{substrate} \quad (2)$$

6.3 Function definition `function_for_Li`

Name function for Li

Arguments kf, Li, Ri, kre, Ci, Ve, Na

Mathematical Expression

$$\frac{kf \cdot Li \cdot Ri - kre \cdot Ci}{Ve \cdot Na} \quad (3)$$

6.4 Function definition `function_for_L`

Name function for L

Arguments kf, L, Rs, kr, Cs, kx, Li, Ve, Na, Y

Mathematical Expression

$$\frac{(kf \cdot L \cdot Rs - kr \cdot Cs - kx \cdot Li \cdot Ve \cdot Na) \cdot Y}{Na} \quad (4)$$

6.5 Function definition `modifier_function_3`

Name modifier function 3

Arguments parameter, modifier

Mathematical Expression

$$parameter \cdot modifier \quad (5)$$

7 Rules

This is an overview of four rules.

7.1 Rule `kf`

Rule kf is an assignment rule for parameter kf:

$$kf = \begin{cases} \frac{kr}{11.1} & \text{if } IL2 = 1 \\ \frac{kr}{8.2} & \text{otherwise} \end{cases} \quad (6)$$

7.2 Rule `kre`

Rule kre is an assignment rule for parameter kre:

$$kre = \begin{cases} kr \cdot 8 & \text{if } IL2 = 1 \\ kr \cdot 5 & \text{otherwise} \end{cases} \quad (7)$$

7.3 Rule kfe

Rule kfe is an assignment rule for parameter kfe :

$$kfe = \begin{cases} \frac{kre}{1000} & \text{if } IL2 = 1 \\ \frac{kre}{3000} & \text{otherwise} \end{cases} \quad (8)$$

7.4 Rule Y_0

Rule Y_0 is a rate rule for species Y_0 :

$$\frac{d}{dt}Y_0 = \begin{cases} \left(\frac{600 \cdot [Cs_0]}{250 + [Cs_0]} - 200 \right) \cdot 1000 & \text{if } \frac{600 \cdot [Cs_0]}{250 + [Cs_0]} - 200 > 0 \\ 0 & \text{otherwise} \end{cases} \quad (9)$$

8 Reactions

This model contains 13 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	Receptor- _binding_to- _Ligand	Receptor binding to Ligand	$Rs_0 \xrightarrow{L_0} Cs_0$	
2	Receptor- _Ligand- _complex- _dissociation	Receptor/Ligand complex dissociation	$Cs_0 \longrightarrow Rs_0$	
3	Upregulation- _of_Receptor- _Synthesis	Upregulation of Receptor Synthesis	$\emptyset \xrightarrow{Cs_0} Rs_0$	
4	Receptor- _Internalisation	Receptor Internalisation	$Rs_0 \longrightarrow Ri_0$	
5	Receptor- _Synthesis	Receptor Synthesis	$\emptyset \longrightarrow Rs_0$	
6	Endocytosis- _of_Receptor- _Ligand_complex	Endocytosis of Receptor/Ligand complex	$Cs_0 \longrightarrow Ci_0$	
7	IL-2_binding- _to_free- _BetaGamma- _heterodimer_1	IL-2 binding to free BetaGamma heterodimer	$Ri_0 \xrightarrow{Li_0} Ci_0$	

Nº	Id	Name	Reaction Equation	SBO
8	Dissocation- _of_IL_2- _Heterodimer- _complex	Dissocation of IL-2/Heterodimer complex	$Ci_0 \longrightarrow Ri_0$	
9	Degradation_of- _heterodimer	Degradation of heterodimer	$Ri_0 \longrightarrow \emptyset$	
10	Degradation- _of_IL_2- _heterodimer- _complex	Degradation of IL-2/heterodimer complex	$Ci_0 \longrightarrow Ld_0$	
11	IL_2_binding- _dissociating	IL-2 binding/dissociating	$Li_0 \xrightarrow{Ri_0, Ci_0} \emptyset$	
12	Ligand- _recycling	Ligand recycling	$Li_0 \longrightarrow \emptyset$	
13	Ligand.Binding- _Dissociation- _Recycling	Ligand Binding/Dissociation/Recycling	$L_0 \xrightarrow{Rs_0, Cs_0, Li_0, Y_0} \emptyset$	

8.1 Reaction Receptor_binding_to_Ligand

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

Name Receptor binding to Ligand

Reaction equation



Reactant

Table 6: Properties of each reactant.

Id	Name	SBO
Rs_0	Rs	

Modifier

Table 7: Properties of each modifier.

Id	Name	SBO
L_0	L	

Product

Table 8: Properties of each product.

Id	Name	SBO
Cs_0	Cs	

Kinetic Law

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

$$v_1 = \text{vol}(\text{COMpartment}) \cdot \text{modifier_function_1}(\text{kf}, [L_0], [Rs_0]) \quad (11)$$

$$\text{modifier_function_1}(\text{parameter}, \text{modifier}, \text{substrate}) = \text{parameter} \cdot \text{modifier} \cdot \text{substrate} \quad (12)$$

$$\text{modifier_function_1}(\text{parameter}, \text{modifier}, \text{substrate}) = \text{parameter} \cdot \text{modifier} \cdot \text{substrate} \quad (13)$$

8.2 Reaction Receptor_Ligand_complex_dissocation

This is an irreversible reaction of one reactant forming one product.

Name Receptor/Ligand complex dissocation

Reaction equation



Reactant

Table 9: Properties of each reactant.

Id	Name	SBO
Cs_0	Cs	

Product

Table 10: Properties of each product.

Id	Name	SBO
Rs_0	Rs	

Kinetic Law

Derived unit $(0.0166667 \text{ s})^{-1} \cdot \text{mol}$

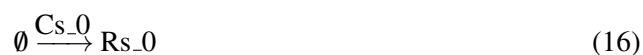
$$v_2 = \text{vol}(\text{COMpartment}) \cdot \text{kr} \cdot [\text{Cs}_0] \quad (15)$$

8.3 Reaction Upregulation_of_Receptor_Synthesis

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

Name Upregulation of Receptor Synthesis

Reaction equation



Modifier

Table 11: Properties of each modifier.

Id	Name	SBO
Cs_0	Cs	

Product

Table 12: Properties of each product.

Id	Name	SBO
Rs_0	Rs	

Kinetic Law

Derived unit $(0.0166667 \text{ s})^{-1} \cdot \text{mol}$

$$v_3 = \text{vol}(\text{COMpartment}) \cdot \text{modifier_function_3}(\text{ksyn}, [\text{Cs}_0]) \quad (17)$$

$$\text{modifier_function_3}(\text{parameter}, \text{modifier}) = \text{parameter} \cdot \text{modifier} \quad (18)$$

$$\text{modifier_function_3}(\text{parameter}, \text{modifier}) = \text{parameter} \cdot \text{modifier} \quad (19)$$

8.4 Reaction `Receptor_Internalisation`

This is an irreversible reaction of one reactant forming one product.

Name Receptor Internalisation

Reaction equation



Reactant

Table 13: Properties of each reactant.

Id	Name	SBO
Rs_0	Rs	

Product

Table 14: Properties of each product.

Id	Name	SBO
Ri_0	Ri	

Kinetic Law

Derived unit $(0.0166667 \text{ s})^{-1} \cdot \text{mol}$

$$v_4 = \text{vol}(\text{COMpartment}) \cdot \text{kt} \cdot [\text{Rs}_0] \quad (21)$$

8.5 Reaction Receptor_Synthesis

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

Name Receptor Synthesis

Reaction equation



Product

Table 15: Properties of each product.

Id	Name	SBO
Rs_0	Rs	

Kinetic Law

Derived unit $1 \cdot (0.0166667 \text{ s})^{-1}$

$$v_5 = \text{vol}(\text{COMpartment}) \cdot \text{Constant_flux_irreversible}(\text{Vs}) \quad (23)$$

$$\text{Constant_flux_irreversible}(v) = v \quad (24)$$

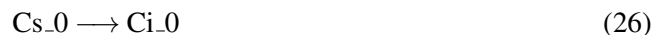
$$\text{Constant_flux_irreversible}(v) = v \quad (25)$$

8.6 Reaction [Endocytosis_of_Receptor_Ligand_complex](#)

This is an irreversible reaction of one reactant forming one product.

Name Endocytosis of Receptor/Ligand complex

Reaction equation



Reactant

Table 16: Properties of each reactant.

Id	Name	SBO
Cs_0	Cs	

Product

Table 17: Properties of each product.

Id	Name	SBO
Ci_0	Ci	

Kinetic Law

Derived unit $(0.0166667 \text{ s})^{-1} \cdot \text{mol}$

$$v_6 = \text{vol}(\text{COMpartment}) \cdot k_e \cdot [\text{Cs}_0] \quad (27)$$

8.7 Reaction [IL_2_binding_to_free_BetaGamma_heterodimer_1](#)

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

Name IL-2 binding to free BetaGamma heterodimer

Reaction equation



Reactant

Table 18: Properties of each reactant.

Id	Name	SBO
Ri_0	Ri	

Modifier

Table 19: Properties of each modifier.

Id	Name	SBO
Li_0	Li	

Product

Table 20: Properties of each product.

Id	Name	SBO
Ci_0	Ci	

Kinetic Law

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

$$v_7 = \text{vol}(\text{COMpartment}) \cdot \text{modifier_function_1}(\text{kfe}, [\text{Li_0}], [\text{Ri_0}]) \quad (29)$$

$$\text{modifier_function_1}(\text{parameter}, \text{modifier}, \text{substrate}) = \text{parameter} \cdot \text{modifier} \cdot \text{substrate} \quad (30)$$

$$\text{modifier_function_1}(\text{parameter}, \text{modifier}, \text{substrate}) = \text{parameter} \cdot \text{modifier} \cdot \text{substrate} \quad (31)$$

8.8 Reaction [Dissocation_of_IL_2_Heterodimer_complex](#)

This is an irreversible reaction of one reactant forming one product.

Name Dissocation of IL-2/Heterodimer complex

Reaction equation



Reactant

Table 21: Properties of each reactant.

Id	Name	SBO
Ci_0	Ci	

Product

Table 22: Properties of each product.

Id	Name	SBO
Ri_0	Ri	

Kinetic Law

Derived unit $(0.0166667 \text{ s})^{-1} \cdot \text{mol}$

$$v_8 = \text{vol}(\text{COMpartment}) \cdot \text{kre} \cdot [\text{Ci}_0] \quad (33)$$

8.9 Reaction Degradation_of_heterodimer

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

Name Degradation of heterodimer

Reaction equation



Reactant

Table 23: Properties of each reactant.

Id	Name	SBO
Ri_0	Ri	

Kinetic Law

Derived unit $(0.0166667 \text{ s})^{-1} \cdot \text{mol}$

$$v_9 = \text{vol}(\text{COMpartment}) \cdot \text{kh} \cdot [\text{Ri}_0] \quad (35)$$

8.10 Reaction Degradation_of_IL_2_heterodimer_complex

This is an irreversible reaction of one reactant forming one product.

Name Degradation of IL-2/heterodimer complex

Reaction equation



Reactant

Table 24: Properties of each reactant.

Id	Name	SBO
Ci_0	Ci	

Product

Table 25: Properties of each product.

Id	Name	SBO
Ld_0	Ld	

Kinetic Law

Derived unit $(0.0166667 \text{ s})^{-1} \cdot \text{mol}$

$$v_{10} = \text{vol}(\text{COMpartment}) \cdot k_h \cdot [\text{Ci}_0] \quad (37)$$

8.11 Reaction IL_2_binding_dissociating

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

Name IL-2 binding/dissociating

Reaction equation



Reactant

Table 26: Properties of each reactant.

Id	Name	SBO
Li_0	Li	

Modifiers

Table 27: Properties of each modifier.

Id	Name	SBO
Ri_0	Ri	
Ci_0	Ci	

Kinetic Law

Derived unit $\text{l}^{-1} \cdot \text{s}^{-1} \cdot (1000000 \text{ mol})^2$

$$v_{11} = \text{vol}(\text{COMpartment}) \cdot \text{function_for_Li}(kf, [\text{Li}_0], [\text{Ri}_0], kre, [\text{Ci}_0], \text{Ve}, \text{NA}) \quad (39)$$

$$\text{function_for_Li}(kf, \text{Li}, \text{Ri}, kre, \text{Ci}, \text{Ve}, \text{Na}) = \frac{kf \cdot \text{Li} \cdot \text{Ri} - kre \cdot \text{Ci}}{\text{Ve} \cdot \text{Na}} \quad (40)$$

$$\text{function_for_Li}(kf, \text{Li}, \text{Ri}, kre, \text{Ci}, \text{Ve}, \text{Na}) = \frac{kf \cdot \text{Li} \cdot \text{Ri} - kre \cdot \text{Ci}}{\text{Ve} \cdot \text{Na}} \quad (41)$$

8.12 Reaction `Ligand_recycling`

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

Name Ligand recycling

Reaction equation



Reactant

Table 28: Properties of each reactant.

Id	Name	SBO
Li_0	Li	

Kinetic Law

Derived unit $(0.0166667 \text{ s})^{-1} \cdot \text{mol}$

$$v_{12} = \text{vol}(\text{COMpartment}) \cdot k_x \cdot [\text{Li}_0] \quad (43)$$

8.13 Reaction `Ligand_Binding_Dissociation_Recycling`

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

Name Ligand Binding/Dissociation/Recycling

Reaction equation



Reactant

Table 29: Properties of each reactant.

Id	Name	SBO
L_0	L	

Modifiers

Table 30: Properties of each modifier.

Id	Name	SBO
Rs_0	Rs	
Cs_0	Cs	
Li_0	Li	
Y_0	Y	

Kinetic Law

Derived unit $\text{l}^{-1} \cdot \text{s}^{-1} \cdot (9999.99999999995 \text{ mol})^3$

$$v_{13} = \text{vol}(\text{COMpartment}) \cdot \text{function_for_L}(\text{kf}, [\text{L}_0], [\text{Rs}_0], \text{kr}, [\text{Cs}_0], \text{kx}, [\text{Li}_0], \text{Ve}, \text{NA}, [\text{Y}_0]) \quad (45)$$

$$\text{function_for_L}(\text{kf}, \text{L}, \text{Rs}, \text{kr}, \text{Cs}, \text{kx}, \text{Li}, \text{Ve}, \text{Na}, \text{Y}) = \frac{(\text{kf} \cdot \text{L} \cdot \text{Rs} - \text{kr} \cdot \text{Cs} - \text{kx} \cdot \text{Li} \cdot \text{Ve} \cdot \text{Na}) \cdot \text{Y}}{\text{Na}} \quad (46)$$

$$\text{function_for_L}(\text{kf}, \text{L}, \text{Rs}, \text{kr}, \text{Cs}, \text{kx}, \text{Li}, \text{Ve}, \text{Na}, \text{Y}) = \frac{(\text{kf} \cdot \text{L} \cdot \text{Rs} - \text{kr} \cdot \text{Cs} - \text{kx} \cdot \text{Li} \cdot \text{Ve} \cdot \text{Na}) \cdot \text{Y}}{\text{Na}} \quad (47)$$

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 `spatialDimensions` > 0 for certain species.

9.1 Species `Rs_0`

Name `Rs`

Notes Initial value of 1500 from paper

Initial concentration $1500 \text{ mol} \cdot \text{l}^{-1}$

This species takes part in six reactions (as a reactant in [Receptor_binding_to_Ligand](#), [Receptor_Internalisation](#) and as a product in [Receptor_Ligand_complex_dissociation](#), [Upregulation_of_Receptor_Synthesis](#), [Receptor_Synthesis](#) and as a modifier in [Ligand_Binding-Dissociation_Recycling](#)).

$$\frac{d}{dt} \text{Rs}_0 = v_2 + v_3 + v_5 - v_1 - v_4 \quad (48)$$

9.2 Species Cs_0

Name Cs

Notes SBML file uploaded with initial value set to 1

Initial concentration 1 mol·l⁻¹

This species takes part in five reactions (as a reactant in [Receptor_Ligand_complex_dissociation](#), [Endocytosis_of_Receptor_Ligand_complex](#) and as a product in [Receptor_binding_to_Ligand](#) and as a modifier in [Upregulation_of_Receptor_Synthesis](#), [Ligand_Binding-Dissociation_Recycling](#)).

$$\frac{d}{dt}Cs_0 = v_1 - v_2 - v_6 \quad (49)$$

9.3 Species Ri_0

Name Ri

Notes Initial value of 300 from paper

Initial concentration 300 mol·l⁻¹

This species takes part in five reactions (as a reactant in [IL_2_binding_to_free_BetaGamma-heterodimer_1](#), [Degradation_of_heterodimer](#) and as a product in [Receptor_Internalisation](#), [Dissociation_of_IL_2_Heterodimer_complex](#) and as a modifier in [IL_2_binding_dissociating](#)).

$$\frac{d}{dt}Ri_0 = v_4 + v_8 - v_7 - v_9 \quad (50)$$

9.4 Species Ci_0

Name Ci

Notes SBML file uploaded with initial value set to 1

Initial concentration 1 mol·l⁻¹

This species takes part in five reactions (as a reactant in [Dissociation_of_IL_2_Heterodimer_complex](#), [Degradation_of_IL_2_heterodimer_complex](#) and as a product in [Endocytosis_of_Receptor_Ligand_complex](#), [IL_2_binding_to_free_BetaGamma_heterodimer_1](#) and as a modifier in [IL_2_binding_dissociating](#)).

$$\frac{d}{dt}Ci_0 = v_6 + v_7 - v_8 - v_{10} \quad (51)$$

9.5 Species Li_0

Name Li

Notes SBML file uploaded with initial value set to 0.01

Initial concentration $0.01 \text{ mol} \cdot \text{l}^{-1}$

This species takes part in four reactions (as a reactant in [IL_2_binding_dissociating](#), [Ligand_recycling](#) and as a modifier in [IL_2_binding_to_free BetaGamma heterodimer_1](#), [Ligand-Binding Dissociation Recycling](#)).

$$\frac{d}{dt}\text{Li}_0 = -v_{11} - v_{12} \quad (52)$$

9.6 Species Ld_0

Name Ld

Notes SBML file uploaded with initial value set to 1

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

This species takes part in one reaction (as a product in [Degradation of IL_2 heterodimer_complex](#)).

$$\frac{d}{dt}\text{Ld}_0 = v_{10} \quad (53)$$

9.7 Species L_0

Name L

Notes SBML file uploaded with initial value set to 10

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

This species takes part in two reactions (as a reactant in [Ligand Binding Dissociation-Recycling](#) and as a modifier in [Receptor binding to Ligand](#)).

$$\frac{d}{dt}\text{L}_0 = -v_{13} \quad (54)$$

9.8 Species Y_0

Name Y

Initial concentration $2.5 \cdot 10^8 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule [Y_0](#)

This species takes part in one reaction (as a modifier in [Ligand Binding Dissociation-Recycling](#)). Not this but one rule determines the species' quantity because this species is on the boundary of the reaction system.

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