SBML Model Report

Model name: "Revilla2003_HIV1therapy"



May 6, 2016

1 General Overview

This is a document in SBML Level 2 Version 4 format. This model was created by the following three authors: Catherine Lloyd¹, Catherine Lloyd² and Catherine Lloyd³ at June 25th 2010 at 1:13 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	1
species types	0	species	0
events	0	constraints	0
reactions	0	function definitions	0
global parameters	16	unit definitions	8
rules	5	initial assignments	0

Model Notes

This a model from the article:

Fighting a virus with a virus: a dynamic model for HIV-1 therapy.

Revilla T, Garcia-Ramos G. Math Biosci 2003 Oct;185(2):191-203 12941536,

Abstract:

¹University of Auckland, c.lloyd@auckland.ac.nz

²University of Auckland, c.lloyd@auckland.ac.nz

³University of Auckland, c.lloyd@auckland.ac.nz

A mathematical model examined a potential therapy for controlling viralinfections using genetically modified viruses. The control of the infection isan indirect effect of the selective elimination by an engineered virus ofinfected cells that are the source of the pathogens. Therefore, this engineered virus could greatly compensate for a dysfunctional immune system compromised by AIDS. In vitro studies using engineered viruses have been shown to decrease the HIV-1 load about 1000-fold. However, the efficacy of this potential treatmentfor reducing the viral load in AIDS patients is unknown. The present modelstudied the interactions among the HIV-1 virus, its main host cell (activated CD4+ T cells), and a therapeutic engineered virus in an in vivo context; and itexamined the conditions for controlling the pathogen. This model predicted asignificant drop in the HIV-1 load, but the treatment does not eradicate HIV. Abasic estimation using a currently engineered virus indicated an HIV-1 loadreduction of 92% and a recovery of host cells to 17% of their normal level. Greater success (98% HIV reduction, 44% host cells recovery) is expected as more competent engineered viruses are designed. These results suggest that therapyusing viruses could be an alternative to extend the survival of AIDS patients.

This model was taken from the CellML repository and automatically converted to SBML. The original model was: **Revilla T, Garcia-Ramos G.** (2003) - version=1.0 The original CellML model was created by:

Catherine Lloyd

c.lloyd@auckland.ac.nz The University of Auckland

This model originates from BioModels Database: A Database of Annotated Published Models (http://www.ebi.ac.uk/biomodels/). It is copyright (c) 2005-2011 The BioModels.net Team. To the extent possible under law, all copyright and related or neighbouring rights to this encoded model have been dedicated to the public domain worldwide. Please refer to CCO Public Domain Dedication for more information.

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

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

2 Unit Definitions

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

2.1 Unit day

Name day

Definition 86400 s

2.2 Unit first_order_rate_constant

Name first_order_rate_constant

Definition $(86400 \text{ s})^{-1}$

2.3 Unit mm3

Name mm3

Definition mm³

2.4 Unit per_mm3

Name per_mm3

Definition mm^{-3}

2.5 Unit cell_per_mm3_day

Name cell_per_mm3_day

Definition $mm^{-3} \cdot (86400 \text{ s})^{-1}$

2.6 Unit vir_per_cell_day

Name vir_per_cell_day

Definition $(86400 \text{ s})^{-1}$

2.7 Unit mm3_per_vir_day

Name mm3_per_vir_day

Definition $mm^3 \cdot (86400 \text{ s})^{-1}$

2.8 Unit time

Name time

Definition 86400 s

2.9 Unit substance

Notes Mole is the predefined SBML unit for substance.

Definition mol

2.10 Unit volume

Notes Litre is the predefined SBML unit for volume.

Definition 1

2.11 Unit area

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

Definition m²

2.12 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			3	1		Ø	

3.1 Compartment COMpartment

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

4 Parameters

This model contains 16 global parameters.

Table 3: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
time-	time		0.000	86400 s	
_environme	11 C				
x	X		3.000		
lamda	lamda		2.000	mm^{-3}	· 🛮
				$(86400 \text{ s})^{-1}$	_

Id	Name	SBO Valu	ue Unit	Constant
d	d	0.0	$010 (86400 \text{ s})^{-1}$	\overline{Z}
У	у	6.0	000	
a	a	0	$330 (86400 \text{ s})^{-1}$	
z	Z	0.0	000	
b	b	2.0	$000 (86400 \text{ s})^{-1}$	\square
v	V	149.	000	
k	k	50.0	$000 (86400 \text{ s})^{-1}$	
u	u	2.0	$000 (86400 \mathrm{s})^{-1}$	\square
W	W	1.	000	
С	c	2000.	,	
q	q	2.0	$000 (86400 \text{ s})^{-1}$	
alpha	alpha	0.0	$1004 \text{mm}^3 \cdot (86400 \text{ s})^{-1}$	
beta	beta	0.0	$004 \text{mm}^3 \cdot (86400 \text{ s})^{-1}$	

5 Rules

This is an overview of five rules.

5.1 Rule x

Rule x is a rate rule for parameter x:

$$\frac{\mathrm{d}}{\mathrm{d}t}\mathbf{x} = 1 \cdot \mathrm{lamda} - (\mathbf{d} \cdot \mathbf{x} + 1 \cdot \mathrm{beta} \cdot \mathbf{x} \cdot \mathbf{v}) \tag{1}$$

5.2 Rule y

Rule y is a rate rule for parameter y:

$$\frac{\mathrm{d}}{\mathrm{d}t}\mathbf{y} = 1 \cdot \text{beta} \cdot \mathbf{x} \cdot \mathbf{v} - (\mathbf{a} \cdot \mathbf{y} + 1 \cdot \text{alpha} \cdot \mathbf{w} \cdot \mathbf{y}) \tag{2}$$

5.3 Rule z

Rule z is a rate rule for parameter z:

$$\frac{\mathrm{d}}{\mathrm{d}t}z = 1 \cdot \mathrm{alpha} \cdot \mathbf{w} \cdot \mathbf{y} - \mathbf{b} \cdot \mathbf{z} \tag{3}$$

5.4 Rule ▽

Rule v is a rate rule for parameter v:

$$\frac{\mathrm{d}}{\mathrm{d}t}\mathbf{v} = \mathbf{k} \cdot \mathbf{y} - \mathbf{u} \cdot \mathbf{v} \tag{4}$$

5.5 Rule ₩

Rule w is a rate rule for parameter w:

$$\frac{\mathrm{d}}{\mathrm{d}t}\mathbf{w} = \mathbf{c} \cdot \mathbf{z} - \mathbf{q} \cdot \mathbf{w} \tag{5}$$

SML2ATEX 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