

SBML Model Report

Model name:
“Maree2006_DuCa_Type1DiabetesModel”



May 6, 2016

1 General Overview

This is a document in SBML Level 2 Version 4 format. This model was created by Ishan Ajmera¹ at October seventh 2011 at 3:36 p. m. and last time modified at October tenth 2014 at 10:36 a. m. Table 1 gives an overview of the quantities of all components of this model.

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

Element	Quantity	Element	Quantity
compartment types	0	compartments	1
species types	0	species	5
events	0	constraints	0
reactions	0	function definitions	0
global parameters	17	unit definitions	0
rules	7	initial assignments	0

Model Notes

This a model from the article:

Modelling the onset of Type 1 diabetes: can impaired macrophage phagocytosis make the difference between health and disease?

Maree AF, Kublik R, Finegood DT, Edelstein-Keshet L. Philos Transact A Math Phys Eng Sci.2006 May 15;364(1842):1267-82. [16608707](#),

¹EMBL-EBI, ajmera@ebi.ac.uk

Abstract:

A wave of apoptosis (programmed cell death) occurs normally in pancreatic beta-cells of newborn mice. We previously showed that macrophages from non-obese diabetic (NOD) mice become activated more slowly and engulf apoptotic cells at a lower rate than macrophages from control (Balb/c) mice. It has been hypothesized that this low clearance could result in secondary necrosis, escalating inflammation and self-antigen presentation that later triggers autoimmune, Type 1 diabetes (T1D). We here investigate whether this hypothesis could offer a reasonable and parsimonious explanation for onset of T1D in NOD mice. We quantify variants of the Copenhagen model (Freiesleben De Blasio et al. 1999 Diabetes 48, 1677), based on parameters from NOD and Balb/c experimental data. We show that the original Copenhagen model fails to explain observed phenomena within a reasonable range of parameter values, predicting an unrealistic all-or-none disease occurrence for both strains. However, if we take into account that, in general, activated macrophages produce harmful cytokines only when engulfing necrotic (but not apoptotic) cells, then the revised model becomes qualitatively and quantitatively reasonable. Further, we show that known differences between NOD and Balb/c mouse macrophage kinetics are large enough to account for the fact that an apoptotic wave can trigger escalating inflammatory response in NOD, but not Balb/c mice. In Balb/c mice, macrophages clear the apoptotic wave so efficiently, that chronic inflammation is prevented.

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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 five unit definitions which are all predefined by SBML and not mentioned in the model.

2.1 Unit substance

Notes Mole is the predefined SBML unit for substance.

Definition mol

2.2 Unit volume

Notes Litre is the predefined SBML unit for volume.

Definition l

2.3 Unit `area`

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

Definition m^2

2.4 Unit `length`

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

Definition m

2.5 Unit `time`

Notes Second is the predefined SBML unit for `time`.

Definition s

3 Compartment

This model contains one compartment.

Table 2: Properties of all compartments.

Id	Name	SBO	Spatial Dimensions	Size	Unit	Constant	Outside
compartment1	compartment1		3	1	litre	<input checked="" type="checkbox"/>	

3.1 Compartment `compartment1`

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

Name `compartment1`

4 Species

This model contains five species. The boundary condition of five of these species is set to `true` so that these species' amount cannot be changed by any reaction. Section 7 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
M	M	compartment1	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
Ma	Ma	compartment1	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
Bn	Bn	compartment1	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
Ba	Ba	compartment1	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
Cy	Cy	compartment1	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>

5 Parameters

This model contains 17 global parameters.

Table 4: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
J	J	0000185	50000.000		<input checked="" type="checkbox"/>
c	c	0000009	0.100		<input checked="" type="checkbox"/>
b	b	0000009	0.090		<input checked="" type="checkbox"/>
d	d	0000009	0.500		<input checked="" type="checkbox"/>
k	k	0000009	0.400		<input checked="" type="checkbox"/>
e1	e1	0000009	10^{-8}		<input checked="" type="checkbox"/>
e2	e2	0000009	10^{-8}		<input checked="" type="checkbox"/>
Amax	Amax	0000009	$2 \cdot 10^7$		<input checked="" type="checkbox"/>
kc	kc	0000009	1.000		<input checked="" type="checkbox"/>
alpha	alpha	0000009	$5 \cdot 10^{-9}$		<input checked="" type="checkbox"/>
delta	delta	0000009	25.000		<input checked="" type="checkbox"/>
g	g	0000009	10^{-5}		<input checked="" type="checkbox"/>
f1	f1	0000009	10^{-5}		<input checked="" type="checkbox"/>
f2	f2	0000009	10^{-5}		<input checked="" type="checkbox"/>
Wmax	Wmax	0000009	$4 \cdot 10^7$		<input checked="" type="checkbox"/>
W	W	0000009	4936.392		<input type="checkbox"/>
parameter_1	x	0000009	9.000		<input type="checkbox"/>

6 Rules

This is an overview of seven rules.

6.1 Rule `parameter_1`

Rule `parameter_1` is an assignment rule for parameter `parameter_1`:

$$\text{parameter_1} = \left(\frac{\text{time} - 9}{3} \right)^2 \quad (1)$$

6.2 Rule `W`

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

$$W = W_{\max} \cdot \exp(\text{parameter_1}) \quad (2)$$

6.3 Rule M

Rule M is a rate rule for species M:

$$\frac{d}{dt}M = J + (k + b) \cdot [Ma] - c \cdot [M] - f1 \cdot [M] \cdot [Ba] - e1 \cdot [M] \cdot ([M] + [Ma]) \quad (3)$$

6.4 Rule Ma

Rule Ma is a rate rule for species Ma:

$$\frac{d}{dt}Ma = f1 \cdot [M] \cdot [Ba] - k \cdot [Ma] - e2 \cdot [Ma] \cdot ([M] + [Ma]) \quad (4)$$

6.5 Rule Bn

Rule Bn is a rate rule for species Bn:

$$\frac{d}{dt}Bn = d \cdot [Ba] - (f1 \cdot [M] + f2 \cdot [Ma]) \cdot [Bn] \quad (5)$$

6.6 Rule Ba

Rule Ba is a rate rule for species Ba:

$$\frac{d}{dt}Ba = W + \frac{Amax \cdot [Cy]}{kc + [Cy]} - (f1 \cdot [M] + f2 \cdot [Ma] + d) \cdot [Ba] \quad (6)$$

6.7 Rule Cy

Rule Cy is a rate rule for species Cy:

$$\frac{d}{dt}Cy = \alpha \cdot [Bn] \cdot [Ma] - \delta \cdot [Cy] \quad (7)$$

7 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.

7.1 Species M

Name M

Initial concentration 477000 mol · l⁻¹

Involved in rule M

One rule determines the species' quantity.

7.2 Species [Ma](#)

Name Ma

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

Involved in rule [Ma](#)

One rule determines the species' quantity.

7.3 Species [Bn](#)

Name Bn

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

Involved in rule [Bn](#)

One rule determines the species' quantity.

7.4 Species [Ba](#)

Name Ba

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

Involved in rule [Ba](#)

One rule determines the species' quantity.

7.5 Species [Cy](#)

Name Cy

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

Involved in rule [Cy](#)

One rule determines the species' quantity.

A Glossary of Systems Biology Ontology Terms

SBO:0000009 kinetic constant: Numerical parameter that quantifies the velocity of a chemical reaction

SBO:0000185 transport reaction: Movement of a physical entity without modification of the structure of the entity

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