

SBML Model Report

Model name: “Cloutier2009 - Brain Energy Metabolism”



May 5, 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, Vijayalakshmi Chelliah² and Audald Lloret i Villas³ at June 25th 2010 at 12:46 a. m. and last time modified at April 17th 2015 at 11:01 a. m. Table 1 shows 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	4
species types	0	species	51
events	0	constraints	0
reactions	0	function definitions	0
global parameters	174	unit definitions	0
rules	100	initial assignments	0

Model Notes

Cloutier2009 - Brain Energy Metabolism

This model was taken from the [CellML repository](#) and automatically converted to SBML. Following

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

²EMBL-EBI, viji@ebi.ac.uk

³EMBL-EBI, lloret@ebi.ac.uk

the submission the parameters are manually encoded and annotated as species and global quantities by BioModels curators.

The original model was: [Cloutier M, Bolger FB, Lowry JP, Wellstead P. \(2009\) -version=1.0](#)

The original CellML model was created by:

Catherine Lloyd

c.lloyd@auckland.ac.nz

The University of Auckland

This model is described in the article: [An integrative dynamic model of brain energy metabolism using in vivo neurochemical measurements](#). Cloutier M, Bolger FB, Lowry JP, Wellstead P. J Comput Neurosci 2009 Dec; 27(3): 391-414

Abstract:

An integrative, systems approach to the modelling of brain energy metabolism is presented. Mechanisms such as glutamate cycling between neurons and astrocytes and glycogen storage in astrocytes have been implemented. A unique feature of the model is its calibration using in vivo data of brain glucose and lactate from freely moving rats under various stimuli. The model has been used to perform simulated perturbation experiments that show that glycogen breakdown in astrocytes is significantly activated during sensory (tail pinch) stimulation. This mechanism provides an additional input of energy substrate during high consumption phases. By way of validation, data from the perfusion of 50 microM propranolol in the rat brain was compared with the model outputs. Propranolol affects the glucose dynamics during stimulation, and this was accurately reproduced in the model by a reduction in the glycogen breakdown in astrocytes. The model's predictive capacity was verified by using data from a sensory stimulation (restraint) that was not used for model calibration. Finally, a sensitivity analysis was conducted on the model parameters, this showed that the control of energy metabolism and transport processes are critical in the metabolic behaviour of cerebral tissue.

This model is hosted on [BioModels Database](#) and identified by: [BIOMD0000000554](#).

To cite BioModels Database, please use: [BioModels Database: An enhanced, curated and annotated resource for published quantitative kinetic models](#).

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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 1

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 Compartments

This model contains four compartments.

Table 2: Properties of all compartments.

Id	Name	SBO	Spatial Dimensions	Size	Unit	Constant	Outside
Astrocytes	Astrocytes		3	1	litre	✓	
Capillaries	Capillaries		3	1	litre	✓	
Extracellular_space	Extracellular space		3	1	litre	✓	
Neurons	Neurons		3	1	litre	✓	

3.1 Compartment Astrocytes

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

Name Astrocytes

Notes Variables indexed $\{\text{straightdbase}\}g\{\text{straightdbase}\}$

3.2 Compartment Capillaries

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

Name Capillaries

Notes Variables indexed $\{\text{\textquotedblbase}c\text{\textquotedblbase}\}$

3.3 Compartment Extracellular_space

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

Name Extracellular space

Notes Variables indexed $\{\text{\textquotedblbase}e\text{\textquotedblbase}\}$

3.4 Compartment Neurons

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

Name Neurons

Notes Variables indexed $\{\text{\textquotedblbase}n\text{\textquotedblbase}\}$

4 Species

This model contains 51 species. The boundary condition of 51 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 Condition
NAg	NAg	Astrocytes	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
GLCg	GLCg	Astrocytes	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
G6Pg	G6Pg	Astrocytes	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
F6Pg	F6Pg	Astrocytes	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
GAPg	GAPg	Astrocytes	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
PEPg	PEPg	Astrocytes	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
PYRg	PYRg	Astrocytes	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
LACg	LACg	Astrocytes	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
NADHg	NADHg	Astrocytes	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
ATPg	ATPg	Astrocytes	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
PCrg	PCrg	Astrocytes	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
O2g	O2g	Astrocytes	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
GLYg	GLYg	Astrocytes	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
GLUg	GLUg	Astrocytes	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
NADg	NADg	Astrocytes	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
ADPg	ADPg	Astrocytes	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
CRg	CRg	Astrocytes	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
NADH_g_tot	NADH_g_tot	Astrocytes	$\text{mol} \cdot \text{l}^{-1}$	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
PCrg_tot	PCrg_tot	Astrocytes	$\text{mol} \cdot \text{l}^{-1}$	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
AMPg	AMPg	Astrocytes	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
O2c	O2c	Capillaries	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>

Id	Name	Compartment	Derived Unit	Constant	Boundary Condition
GLCc	GLCc	Capillaries	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
LACc	LACc	Capillaries	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
CO2c	CO2c	Capillaries	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
GLCe	GLCe	Extracellular_space	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
LACe	LACe	Extracellular_space	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
GLUe	GLUe	Extracellular_space	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
NAe	NAe	Extracellular_space	$\text{mol} \cdot \text{l}^{-1}$	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
O2a	O2a	Extracellular_space	$\text{mol} \cdot \text{l}^{-1}$	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
CO2a	CO2a	Extracellular_space	$\text{mol} \cdot \text{l}^{-1}$	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
GLCa	GLCa	Extracellular_space	$\text{mol} \cdot \text{l}^{-1}$	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
LACa	LACa	Extracellular_space	$\text{mol} \cdot \text{l}^{-1}$	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
NAn	NAn	Neurons	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
GLCn	GLCn	Neurons	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
G6Pn	G6Pn	Neurons	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
F6Pn	F6Pn	Neurons	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
GAPn	GAPn	Neurons	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
PEPn	PEPn	Neurons	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
PYRn	PYRn	Neurons	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
LACn	LACn	Neurons	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
NADHn	NADHn	Neurons	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
ATPn	ATPn	Neurons	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
PCrn	PCrn	Neurons	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
O2n	O2n	Neurons	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
GLUn	GLUn	Neurons	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
ADPn	ADPn	Neurons	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
CRn	CRn	Neurons	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>
NADn	NADn	Neurons	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>

Id	Name	Compartment	Derived Unit	Constant	Boundary Condi- tion
NADH_n_tot	NADH_n_tot	Neurons	$\text{mol} \cdot \text{l}^{-1}$	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
PCrn_tot	PCrn_tot	Neurons	$\text{mol} \cdot \text{l}^{-1}$	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
AMPn	AMPn	Neurons	$\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input checked="" type="checkbox"/>

5 Parameters

This model contains 174 global parameters.

Table 4: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
u_n	u_n		1.042		<input type="checkbox"/>
u_g	u_g		1.075		<input type="checkbox"/>
dAMP_dATPn	dAMP_dATPn		−0.101		<input type="checkbox"/>
dAMP_dATPg	dAMP_dATPg		−0.116		<input type="checkbox"/>
Vv	Vv		0.024		<input type="checkbox"/>
dHb	dHb		0.022		<input type="checkbox"/>
Vn_leak_Na	Vn_leak_Na		0.475		<input type="checkbox"/>
gn_NA	gn_NA		0.004		<input checked="" type="checkbox"/>
Vn_pump	Vn_pump		0.158		<input type="checkbox"/>
Vn_stim	Vn_stim		0.000		<input type="checkbox"/>
V_en_GLC	V_en_GLC		0.006		<input type="checkbox"/>
Km_en_GLC	Km_en_GLC		5.320		<input checked="" type="checkbox"/>
Vm_en_GLC	Vm_en_GLC		0.504		<input checked="" type="checkbox"/>
Vn_hk	Vn_hk		0.006		<input type="checkbox"/>
Vmax_n_hk	Vmax_n_hk		0.051		<input checked="" type="checkbox"/>
Vn_pgi	Vn_pgi		0.006		<input type="checkbox"/>
Vmaxf_n_pgi	Vmaxf_n_pgi		0.500		<input checked="" type="checkbox"/>
Vmaxr_n_pgi	Vmaxr_n_pgi		0.450		<input checked="" type="checkbox"/>
Vn_pfk	Vn_pfk		0.006		<input type="checkbox"/>
kn_pfk	kn_pfk		0.558		<input checked="" type="checkbox"/>
Vn_pgk	Vn_pgk		0.012		<input type="checkbox"/>
kn_pgk	kn_pgk		0.429		<input checked="" type="checkbox"/>
Vn_pk	Vn_pk		0.012		<input type="checkbox"/>
kn_pk	kn_pk		28.600		<input checked="" type="checkbox"/>
Vn_ldh	Vn_ldh		−0.001		<input type="checkbox"/>
kfn_ldh	kfn_ldh		5.300		<input checked="" type="checkbox"/>
kfn_ldh	kfn_ldh		0.105		<input checked="" type="checkbox"/>
Vn_mito	Vn_mito		0.013		<input type="checkbox"/>
Vmax_n_mito	Vmax_n_mito		0.056		<input checked="" type="checkbox"/>
Vne_LAC	Vne_LAC		−0.001		<input type="checkbox"/>
Vmax_ne_LAC	Vmax_ne_LAC		0.198		<input checked="" type="checkbox"/>
Km_ne_LAC	Km_ne_LAC		0.093		<input checked="" type="checkbox"/>
Vn_ATPase	Vn_ATPase		0.049		<input type="checkbox"/>
Vmax_n-ATPase	Vmax_n-ATPase		0.049		<input checked="" type="checkbox"/>
Vn_ck	Vn_ck		$2.93701651940294 \cdot 10^{-5}$		<input type="checkbox"/>
kfn_ck	kfn_ck		0.015		<input checked="" type="checkbox"/>

Id	Name	SBO	Value	Unit	Constant
kfn_ck	kfn_ck		0.052		<input checked="" type="checkbox"/>
Vcn_02	Vcn_02		0.039		<input type="checkbox"/>
nh_02.Vcn_02	nh_02		2.700		<input checked="" type="checkbox"/>
PScapn	PScapn		0.220		<input checked="" type="checkbox"/>
Vg_leak_Na	Vg_leak_Na		0.190		<input type="checkbox"/>
gg_NA	gg_NA		0.003		<input checked="" type="checkbox"/>
Vg_pump	Vg_pump		0.063		<input type="checkbox"/>
Veg_GLC	Veg_GLC		0.002		<input type="checkbox"/>
Km_eg_GLC	Km_eg_GLC		3.530		<input checked="" type="checkbox"/>
Vm_eg_GLC	Vm_eg_GLC		0.038		<input checked="" type="checkbox"/>
Vcg_GLC	Vcg_GLC		0.003		<input type="checkbox"/>
Km_cg_GLC	Km_cg_GLC		9.920		<input checked="" type="checkbox"/>
Vm_cg_GLC	Vm_cg_GLC		0.010		<input checked="" type="checkbox"/>
Vg_hk	Vg_hk		0.005		<input type="checkbox"/>
Vmax_g_hk	Vmax_g_hk		0.050		<input checked="" type="checkbox"/>
Vg_pgi	Vg_pgi		0.005		<input type="checkbox"/>
Vmaxf_g_pgi	Vmaxf_g_pgi		0.500		<input checked="" type="checkbox"/>
Vmaxr_g_pgi	Vmaxr_g_pgi		0.450		<input checked="" type="checkbox"/>
Vg_pfk	Vg_pfk		0.005		<input type="checkbox"/>
kg_pfk	kg_pfk		0.403		<input checked="" type="checkbox"/>
Vg_pgk	Vg_pgk		0.009		<input type="checkbox"/>
kg_pgk	kg_pgk		0.251		<input checked="" type="checkbox"/>
Vg_pk	Vg_pk		0.009		<input type="checkbox"/>
kg_pk	kg_pk		2.730		<input checked="" type="checkbox"/>
Vg_ldh	Vg_ldh		0.003		<input type="checkbox"/>
kfg_ldh	kfg_ldh		6.261		<input checked="" type="checkbox"/>
krp_ldh	krp_ldh		0.547		<input checked="" type="checkbox"/>
Vg_mito	Vg_mito		0.006		<input type="checkbox"/>
Vmax_g_mito	Vmax_g_mito		0.008		<input checked="" type="checkbox"/>
Vge_LAC	Vge_LAC		0.003		<input type="checkbox"/>
Vmax_ge_LAC	Vmax_ge_LAC		0.086		<input checked="" type="checkbox"/>
Km_ge_LAC	Km_ge_LAC		0.222		<input checked="" type="checkbox"/>
Vgc_LAC	Vgc_LAC		$1.46095762940601 \cdot 10^{-5}$		<input type="checkbox"/>
Vmax_gc_LAC	Vmax_gc_LAC		$2.1856 \cdot 10^{-4}$		<input checked="" type="checkbox"/>
Km_gc_LAC	Km_gc_LAC		0.129		<input checked="" type="checkbox"/>
Vg_ATPase	Vg_ATPase		0.036		<input type="checkbox"/>
Vmax_g-ATPase	Vmax_g-ATPase		0.036		<input checked="" type="checkbox"/>
Vg_ck	Vg_ck		$8.98869880248884 \cdot 10^{-5}$		<input type="checkbox"/>
krp_ck	krp_ck		0.021		<input checked="" type="checkbox"/>
kfg_ck	kfg_ck		0.024		<input checked="" type="checkbox"/>
Vcg_02	Vcg_02		0.018		<input type="checkbox"/>

Id	Name	SBO	Value	Unit	Constant
PScapg	PScapg		0.246		<input checked="" type="checkbox"/>
Vc_O2	Vc_O2		4.014		<input type="checkbox"/>
Vc_GLC	Vc_GLC		0.698		<input type="checkbox"/>
Vce_GLC	Vce_GLC		0.015		<input type="checkbox"/>
Km_ce_GLC	Km_ce_GLC		8.457		<input checked="" type="checkbox"/>
Vm_ce_GLC	Vm_ce_GLC		0.049		<input checked="" type="checkbox"/>
Vc_LAC	Vc_LAC		-0.053		<input type="checkbox"/>
Vec_LAC	Vec_LAC		0.001		<input type="checkbox"/>
Km_ec_LAC	Km_ec_LAC		0.765		<input checked="" type="checkbox"/>
Vm_ec_LAC	Vm_ec_LAC		0.033		<input checked="" type="checkbox"/>
Vnc_CO2	Vnc_CO2		0.039		<input type="checkbox"/>
Vgc_CO2	Vgc_CO2		0.018		<input type="checkbox"/>
Vn_stim_GLU	Vn_stim_GLU		0.000		<input type="checkbox"/>
Vg_gs	Vg_gs		0.000		<input type="checkbox"/>
Vmax_g_gs	Vmax_g_gs		0.300		<input checked="" type="checkbox"/>
Veg_GLU	Veg_GLU		0.000		<input type="checkbox"/>
Vmax_eg_GLU	Vmax_eg_GLU		0.021		<input checked="" type="checkbox"/>
Vc_CO2	Vc_CO2		4.015		<input type="checkbox"/>
Vg_glys	Vg_glys	9.08171994158688 · 10 ⁻⁵			<input type="checkbox"/>
Vmax_glys	Vmax_glys		1.528 · 10 ⁻⁴		<input checked="" type="checkbox"/>
Km_G6P_glys	Km_G6P_glys		0.500		<input checked="" type="checkbox"/>
Vg_glyp	Vg_glyp	3.51571428571429 · 10 ⁻⁵			<input type="checkbox"/>
Vmax_glyp	Vmax_glyp		4.922 · 10 ⁻⁵		<input checked="" type="checkbox"/>
Km_GLY	Km_GLY		1.000		<input checked="" type="checkbox"/>
deltaVt_GLY	deltaVt_GLY		1.000		<input type="checkbox"/>
Fin_t	Fin_t		0.012		<input type="checkbox"/>
CBF0_Fin_t	CBF0		0.012		<input checked="" type="checkbox"/>
Fout_t	Fout_t		0.012		<input type="checkbox"/>
BOLD	BOLD		0.042		<input type="checkbox"/>
v_stim	v_stim		0.000		<input type="checkbox"/>
unitpulseSB	unitpulseSB		0.000		<input type="checkbox"/>
unitstepSB	unitstepSB		0.000		<input type="checkbox"/>
Km_PYR	Km_PYR		0.063		<input checked="" type="checkbox"/>
Km_ATP	Km_ATP		0.015		<input checked="" type="checkbox"/>
Ki_ATP	Ki_ATP		0.760		<input checked="" type="checkbox"/>
Km_ADP	Km_ADP		0.001		<input checked="" type="checkbox"/>
Km_O2	Km_O2		0.003		<input checked="" type="checkbox"/>
Km_GLC	Km_GLC		0.105		<input checked="" type="checkbox"/>
Km_GLU	Km_GLU		0.050		<input checked="" type="checkbox"/>
Km_G6P	Km_G6P		0.500		<input checked="" type="checkbox"/>
Km_F6P_pgi	Km_F6P_pgi		0.060		<input checked="" type="checkbox"/>
Km_F6P_pfk	Km_F6P_pfk		0.180		<input checked="" type="checkbox"/>

Id	Name	SBO	Value	Unit	Constant
Km_pump	Km_pump		0.424		<input checked="" type="checkbox"/>
nh_O2_model- _parameters	nh_O2_2		2.700		<input checked="" type="checkbox"/>
Ko2	Ko2		0.090		<input checked="" type="checkbox"/>
kpump	kpump		$3.17 \cdot 10^{-7}$		<input checked="" type="checkbox"/>
ATPtot	ATPtot		2.379		<input checked="" type="checkbox"/>
nH	nH		4.000		<input checked="" type="checkbox"/>
nOP	nOP		15.000		<input checked="" type="checkbox"/>
NAero	NAero		3.000		<input checked="" type="checkbox"/>
Rng	Rng		1.800		<input checked="" type="checkbox"/>
Reg	Reg		0.800		<input checked="" type="checkbox"/>
Ren	Ren		0.444		<input checked="" type="checkbox"/>
Rcn	Rcn		0.012		<input checked="" type="checkbox"/>
Rcg	Rcg		0.022		<input checked="" type="checkbox"/>
Rce	Rce		0.028		<input checked="" type="checkbox"/>
Sm_n	Sm_n		40500.000		<input checked="" type="checkbox"/>
Vm	Vm		-70.000		<input checked="" type="checkbox"/>
RT	RT		2577340.000		<input checked="" type="checkbox"/>
F	F		96500.000		<input checked="" type="checkbox"/>
Vn	Vn		0.450		<input checked="" type="checkbox"/>
G6P_inh_hk	G6P_inh_hk		0.600		<input checked="" type="checkbox"/>
aG6P_inh_hk	aG6P_inh_hk		20.000		<input checked="" type="checkbox"/>
rATP_mito	rATP_mito		20.000		<input checked="" type="checkbox"/>
aATP_mito	aATP_mito		5.000		<input checked="" type="checkbox"/>
HbOP	HbOP		8.600		<input checked="" type="checkbox"/>
Sm_g	Sm_g		10500.000		<input checked="" type="checkbox"/>
Vg	Vg		0.250		<input checked="" type="checkbox"/>
KO1	KO1		1.000		<input checked="" type="checkbox"/>
Vc	Vc		0.006		<input checked="" type="checkbox"/>
R_GLU_NA	R_GLU_NA		0.075		<input checked="" type="checkbox"/>
KO2	KO2		1.000		<input checked="" type="checkbox"/>
KO3	KO3		1.000		<input checked="" type="checkbox"/>
GLY_inh	GLY_inh		4.200		<input checked="" type="checkbox"/>
aGLY_inh	aGLY_inh		20.000		<input checked="" type="checkbox"/>
CBF0_model- _parameters	CBF0_2		0.012		<input checked="" type="checkbox"/>
Vv0	Vv0		0.024		<input checked="" type="checkbox"/>
tv	tv		35.000		<input checked="" type="checkbox"/>
qak	qak		0.920		<input checked="" type="checkbox"/>
k1	k1		2.220		<input checked="" type="checkbox"/>
k2	k2		0.460		<input checked="" type="checkbox"/>
k3	k3		0.430		<input checked="" type="checkbox"/>

Id	Name	SBO	Value	Unit	Constant
dHb0	dHb0		0.064		<input checked="" type="checkbox"/>
stim	stim		1.000		<input checked="" type="checkbox"/>
to	to		200.000		<input checked="" type="checkbox"/>
tend	tend		300.000		<input checked="" type="checkbox"/>
v1_n	v1_n		0.041		<input checked="" type="checkbox"/>
v2_n	v2_n		2.550		<input checked="" type="checkbox"/>
t_n_stim	t_n_stim		2.000		<input checked="" type="checkbox"/>
sr	sr		4.592		<input checked="" type="checkbox"/>
t1	t1		2.000		<input checked="" type="checkbox"/>
delta_GLY	delta_GLY		62.000		<input checked="" type="checkbox"/>
deltaf	deltaf		0.420		<input checked="" type="checkbox"/>
tend_GLY	tend_GLY		440.000		<input checked="" type="checkbox"/>
to_GLY	to_GLY		83.000		<input checked="" type="checkbox"/>
sr_GLY	sr_GLY		4.000		<input checked="" type="checkbox"/>
unitstepSB2	unitstepSB2		0.000		<input type="checkbox"/>

6 Rules

This is an overview of 100 rules.

6.1 Rule V_{en_GLC}

Rule V_{en_GLC} is an assignment rule for parameter V_{en_GLC} :

$$V_{en_GLC} = V_{m_en_GLC} \cdot \left(\frac{[GLCe]}{[GLCe] + K_{m_en_GLC}} - \frac{[GLCn]}{[GLCn] + K_{m_en_GLC}} \right) \quad (1)$$

6.2 Rule V_{n_hk}

Rule V_{n_hk} is an assignment rule for parameter V_{n_hk} :

$$V_{n_hk} = V_{max_n_hk} \cdot [ATPn] \cdot \frac{[GLCn]}{[GLCn] + K_{m_GLC}} \cdot \left(1 - \frac{1}{1 + \exp(a_{G6P_inh_hk} \cdot 1 \cdot ([G6Pn] - G6P_{inh_hk}))} \right) \quad (2)$$

6.3 Rule V_{n_pgi}

Rule V_{n_pgi} is an assignment rule for parameter V_{n_pgi} :

$$V_{n_pgi} = V_{maxf_n_pgi} \cdot \frac{[G6Pn]}{[G6Pn] + K_{m_G6P}} - V_{maxr_n_pgi} \cdot \frac{[F6Pn]}{[F6Pn] + K_{m_F6P_pgi}} \quad (3)$$

6.4 Rule NADg

Rule NADg is an assignment rule for species NADg:

$$\text{NADg} = [\text{NADH_g_tot}] - [\text{NADHg}] \quad (4)$$

Derived unit mol · l⁻¹

6.5 Rule ADPg

Rule ADPg is an assignment rule for species ADPg:

$$\text{ADPg} = \frac{[\text{ATPg}]}{2} \cdot \left(\text{qak} + \left(\text{qak}^2 + 4 \cdot \text{qak} \cdot \left(\frac{\text{ATPtot}}{[\text{ATPg}]} - 1 \right) \right)^{\frac{1}{2}} \right) \quad (5)$$

6.6 Rule CRg

Rule CRg is an assignment rule for species CRg:

$$\text{CRg} = [\text{PCrg_tot}] - [\text{PCrg}] \quad (6)$$

Derived unit mol · l⁻¹

6.7 Rule AMPg

Rule AMPg is an assignment rule for species AMPg:

$$\text{AMPg} = \text{ATPtot} - ([\text{ATPg}] + [\text{ADPg}]) \quad (7)$$

6.8 Rule Vg_ck

Rule Vg_ck is an assignment rule for parameter Vg_ck:

$$\text{Vg_ck} = \text{kfg_ck} \cdot [\text{PCrg}] \cdot [\text{ADPg}] - \text{krg_ck} \cdot [\text{CRg}] \cdot [\text{ATPg}] \quad (8)$$

6.9 Rule ADPn

Rule ADPn is an assignment rule for species ADPn:

$$\text{ADPn} = \frac{[\text{ATPn}]}{2} \cdot \left(\text{qak} + \left(\text{qak}^2 + 4 \cdot \text{qak} \cdot \left(\frac{\text{ATPtot}}{[\text{ATPn}]} - 1 \right) \right)^{\frac{1}{2}} \right) \quad (9)$$

6.10 Rule CRn

Rule CRn is an assignment rule for species CRn:

$$\text{CRn} = [\text{PCrn_tot}] - [\text{PCrn}] \quad (10)$$

Derived unit mol · l⁻¹

6.11 Rule NADn

Rule NADn is an assignment rule for species NADn :

$$\text{NADn} = [\text{NADH.n.tot}] - [\text{NADHn}] \quad (11)$$

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

6.12 Rule AMPn

Rule AMPn is an assignment rule for species AMPn :

$$\text{AMPn} = \text{ATPtot} - ([\text{ATPn}] + [\text{ADPn}]) \quad (12)$$

6.13 Rule u.n

Rule u.n is an assignment rule for parameter u.n :

$$\text{u.n} = \text{qak}^2 + 4 \cdot \text{qak} \cdot \left(\frac{\text{ATPtot}}{[\text{ATPn}]} - 1 \right) \quad (13)$$

6.14 Rule u.g

Rule u.g is an assignment rule for parameter u.g :

$$\text{u.g} = \text{qak}^2 + 4 \cdot \text{qak} \cdot \left(\frac{\text{ATPtot}}{[\text{ATPg}]} - 1 \right) \quad (14)$$

6.15 Rule dAMP.dATPn

Rule dAMP.dATPn is an assignment rule for parameter dAMP.dATPn :

$$\text{dAMP.dATPn} = \frac{\text{qak}}{2} + \text{qak} \cdot \frac{\text{ATPtot}}{[\text{ATPn}] \cdot \text{u.n}^{\frac{1}{2}}} - \left(1 + 0.5 \cdot \text{u.n}^{\frac{1}{2}} \right) \quad (15)$$

6.16 Rule dAMP.dATPg

Rule dAMP.dATPg is an assignment rule for parameter dAMP.dATPg :

$$\text{dAMP.dATPg} = \frac{\text{qak}}{2} + \text{qak} \cdot \frac{\text{ATPtot}}{[\text{ATPg}] \cdot \text{u.g}^{\frac{1}{2}}} - \left(1 + 0.5 \cdot \text{u.g}^{\frac{1}{2}} \right) \quad (16)$$

6.17 Rule Vn.leak.Na

Rule Vn.leak.Na is an assignment rule for parameter Vn.leak.Na :

$$\text{Vn.leak.Na} = \frac{\text{Sm.n}}{\text{Vn}} \cdot \frac{\text{gn.Na}}{\text{F}} \cdot \left(\frac{\text{RT}}{\text{F}} \cdot \left(\frac{[\text{NAe}]}{[\text{NAn}]} \right) - \text{Vm} \right) \quad (17)$$

6.18 Rule Vn_pump

Rule Vn_pump is an assignment rule for parameter Vn_pump :

$$Vn_pump = \frac{Sm_n}{Vn} \cdot kpump \cdot [ATPn] \cdot [NAn] \cdot \left(1 + \frac{[ATPn]}{Km_pump}\right)^1 \quad (18)$$

6.19 Rule Vn_pfk

Rule Vn_pfk is an assignment rule for parameter Vn_pfk :

$$Vn_pfk = kn_pfk \cdot [ATPn] \cdot \frac{[F6Pn]}{[F6Pn] + Km_F6P_pfk} \cdot \left(1 + \left(\frac{[ATPn]}{Ki_ATP}\right)^{nH}\right)^1 \quad (19)$$

6.20 Rule Vn_pgk

Rule Vn_pgk is an assignment rule for parameter Vn_pgk :

$$Vn_pgk = kn_pgk \cdot [GAPn] \cdot [ADPn] \cdot \frac{[NADn]}{[NADHn]} \quad (20)$$

6.21 Rule Vn_pk

Rule Vn_pk is an assignment rule for parameter Vn_pk :

$$Vn_pk = kn_pk \cdot [PEPn] \cdot [ADPn] \quad (21)$$

6.22 Rule Vn_ldh

Rule Vn_ldh is an assignment rule for parameter Vn_ldh :

$$Vn_ldh = kfn_ldh \cdot [PYRn] \cdot [NADHn] - krm_ldh \cdot [LACn] \cdot [NADn] \quad (22)$$

6.23 Rule Vn_mito

Rule Vn_mito is an assignment rule for parameter Vn_mito :

$$Vn_mito = Vmax_n_mito \cdot \frac{[O2n]}{[O2n] + Km_O2} \cdot \frac{[ADPn]}{[ADPn] + Km_ADP} \cdot \frac{[PYRn]}{[PYRn] + Km_PYR} \cdot \left(1 - \frac{1}{1 + \exp\left(aATP_mito \cdot 1 \cdot \left(\frac{[ATPn]}{[ADPn]} - 1 \cdot rATP_mito\right)\right)}\right) \quad (23)$$

6.24 Rule Vne_LAC

Rule Vne_LAC is an assignment rule for parameter Vne_LAC :

$$Vne_LAC = Vmax_ne_LAC \cdot \left(\frac{[LACn]}{[LACn] + Km_ne_LAC} - \frac{[LACe]}{[LACe] + Km_ne_LAC}\right) \quad (24)$$

6.25 Rule Vn_ATPase

Rule Vn_ATPase is an assignment rule for parameter Vn_ATPase :

$$Vn_ATPase = Vmax_n_ATPase \cdot \frac{[ATPn]}{[ATPn] + 0.0010} \quad (25)$$

6.26 Rule Vge_LAC

Rule Vge_LAC is an assignment rule for parameter Vge_LAC :

$$Vge_LAC = Vmax_ge_LAC \cdot \left(\frac{[LACg]}{[LACg] + Km_ge_LAC} - \frac{[LACe]}{[LACe] + Km_ge_LAC} \right) \quad (26)$$

6.27 Rule Vn_ck

Rule Vn_ck is an assignment rule for parameter Vn_ck :

$$Vn_ck = kfn_ck \cdot [PCrn] \cdot [ADPn] - kmn_ck \cdot [CRn] \cdot [ATPn] \quad (27)$$

6.28 Rule Vcn_O2

Rule Vcn_O2 is an assignment rule for parameter Vcn_O2 :

$$Vcn_O2 = \frac{PScapn}{Vn} \cdot \left(Ko2 \cdot \left(\frac{HbOP}{[O2c]} - 1 \right)^{\frac{1}{nh.O2.Vcn.O2}} - [O2n] \right) \quad (28)$$

6.29 Rule Vg_leak_Na

Rule Vg_leak_Na is an assignment rule for parameter Vg_leak_Na :

$$Vg_leak_Na = \frac{Sm_g}{Vg} \cdot \frac{gg_NA}{F} \cdot \left(\frac{RT}{F} \cdot \left(\frac{[NAe]}{[NAg]} \right) - Vm \right) \quad (29)$$

6.30 Rule Vg_pump

Rule Vg_pump is an assignment rule for parameter Vg_pump :

$$Vg_pump = \frac{Sm_g}{Vg} \cdot kpump \cdot [ATPg] \cdot [NAg] \cdot \left(1 + \frac{[ATPg]}{Km_pump} \right)^1 \quad (30)$$

6.31 Rule Veg_GLC

Rule Veg_GLC is an assignment rule for parameter Veg_GLC :

$$Veg_GLC = KO1 \cdot Vm_eg_GLC \cdot \left(\frac{[GLCe]}{[GLCe] + Km_eg_GLC} - \frac{[GLCg]}{[GLCg] + Km_eg_GLC} \right) \quad (31)$$

6.32 Rule V_{cg_GLC}

Rule V_{cg_GLC} is an assignment rule for parameter V_{cg_GLC} :

$$V_{cg_GLC} = V_{m_cg_GLC} \cdot \left(\frac{[GLCc]}{[GLCc] + K_{m_cg_GLC}} - \frac{[GLCg]}{[GLCg] + K_{m_cg_GLC}} \right) \quad (32)$$

6.33 Rule V_{g_hk}

Rule V_{g_hk} is an assignment rule for parameter V_{g_hk} :

$$V_{g_hk} = V_{max_g_hk} \cdot [ATPg] \cdot \frac{[GLCg]}{[GLCg] + K_{m_GLC}} \cdot \left(1 - \frac{1}{1 + \exp(a_{G6P_inh_hk} \cdot 1 \cdot ([G6Pg] - G6P_inh_hk))} \right) \quad (33)$$

6.34 Rule V_{g_pgi}

Rule V_{g_pgi} is an assignment rule for parameter V_{g_pgi} :

$$V_{g_pgi} = V_{maxf_g_pgi} \cdot \frac{[G6Pg]}{[G6Pg] + K_{m_G6P}} - V_{maxr_g_pgi} \cdot \frac{[F6Pg]}{[F6Pg] + K_{m_F6P_pgi}} \quad (34)$$

6.35 Rule V_{g_pfk}

Rule V_{g_pfk} is an assignment rule for parameter V_{g_pfk} :

$$V_{g_pfk} = k_{g_pfk} \cdot [ATPg] \cdot \frac{[F6Pg]}{[F6Pg] + K_{m_F6P_pfk}} \cdot \left(1 + \left(\frac{[ATPg]}{K_{i_ATP}} \right)^{nH} \right)^1 \quad (35)$$

6.36 Rule V_{g_pgk}

Rule V_{g_pgk} is an assignment rule for parameter V_{g_pgk} :

$$V_{g_pgk} = k_{g_pgk} \cdot [GAPg] \cdot [ADPg] \cdot \frac{[NADg]}{[NADHg]} \quad (36)$$

6.37 Rule V_{g_pk}

Rule V_{g_pk} is an assignment rule for parameter V_{g_pk} :

$$V_{g_pk} = k_{g_pk} \cdot [PEPg] \cdot [ADPg] \quad (37)$$

6.38 Rule V_{g_ldh}

Rule V_{g_ldh} is an assignment rule for parameter V_{g_ldh} :

$$V_{g_ldh} = k_{fg_ldh} \cdot [PYRg] \cdot [NADHg] - k_{rg_ldh} \cdot [LACg] \cdot [NADg] \quad (38)$$

6.39 Rule Vg_mito

Rule Vg_mito is an assignment rule for parameter Vg_mito :

$$Vg_mito = Vmax_g_mito \cdot \frac{[O2g]}{[O2g] + Km_O2} \cdot \frac{[ADPg]}{[ADPg] + Km_ADP} \cdot \frac{[PYRg]}{[PYRg] + Km_PYR} \cdot \left(1 - \frac{1}{1 + \exp\left(1 \cdot (aATP_mito) \cdot \left(\frac{[ATPg]}{[ADPg]} - 1 \cdot rATP_mito\right)\right)} \right) \quad (39)$$

6.40 Rule Vgc_LAC

Rule Vgc_LAC is an assignment rule for parameter Vgc_LAC :

$$Vgc_LAC = Vmax_gc_LAC \cdot \left(\frac{[LACg]}{[LACg] + Km_gc_LAC} - \frac{[LACc]}{[LACc] + Km_gc_LAC} \right) \quad (40)$$

6.41 Rule Vg_ATPase

Rule Vg_ATPase is an assignment rule for parameter Vg_ATPase :

$$Vg_ATPase = Vmax_g_ATPase \cdot \frac{[ATPg]}{[ATPg] + 0.0010} \quad (41)$$

6.42 Rule Vcg_O2

Rule Vcg_O2 is an assignment rule for parameter Vcg_O2 :

$$Vcg_O2 = \frac{PScapg}{Vg} \cdot \left(Ko2 \cdot \left(\frac{HbOP}{[O2c]} - 1 \right)^{\frac{1}{nh.O2.model.parameters}} - [O2g] \right) \quad (42)$$

6.43 Rule Vce_GLC

Rule Vce_GLC is an assignment rule for parameter Vce_GLC :

$$Vce_GLC = Vm_ce_GLC \cdot \left(\frac{[GLCc]}{[GLCc] + Km_ce_GLC} - \frac{[GLCe]}{[GLCe] + Km_ce_GLC} \right) \quad (43)$$

6.44 Rule Vec_LAC

Rule Vec_LAC is an assignment rule for parameter Vec_LAC :

$$Vec_LAC = Vm_ec_LAC \cdot \left(\frac{[LACe]}{[LACe] + Km_ec_LAC} - \frac{[LACc]}{[LACc] + Km_ec_LAC} \right) \quad (44)$$

6.45 Rule V_{nc_CO2}

Rule V_{nc_CO2} is an assignment rule for parameter V_{nc_CO2} :

$$V_{nc_CO2} = 3 \cdot V_{n_mito} \quad (45)$$

6.46 Rule V_{gc_CO2}

Rule V_{gc_CO2} is an assignment rule for parameter V_{gc_CO2} :

$$V_{gc_CO2} = 3 \cdot V_{g_mito} \quad (46)$$

6.47 Rule V_{g_gs}

Rule V_{g_gs} is an assignment rule for parameter V_{g_gs} :

$$V_{g_gs} = V_{max_g_gs} \cdot \frac{[GLUg]}{[GLUg] + K_{m_GLU}} \cdot \frac{[ATPg]}{[ATPg] + K_{m_ATP}} \quad (47)$$

6.48 Rule V_{eg_GLU}

Rule V_{eg_GLU} is an assignment rule for parameter V_{eg_GLU} :

$$V_{eg_GLU} = V_{max_eg_GLU} \cdot \frac{[GLUe]}{[GLUe] + K_{m_GLU}} \quad (48)$$

6.49 Rule V_{g_glys}

Rule V_{g_glys} is an assignment rule for parameter V_{g_glys} :

$$V_{g_glys} = V_{max_glys} \cdot \frac{[G6Pg]}{[G6Pg] + K_{m_G6P_glys}} \cdot \left(1 - \frac{1}{1 + \exp(a_{GLY_inh} \cdot 1 \cdot ([GLYg] - GLY_inh))} \right) \quad (49)$$

6.50 Rule Fin_t

Rule Fin_t is an assignment rule for parameter Fin_t :

$$Fin_t = CBF0_Fin_t + \left(stim \cdot CBF0_Fin_t \cdot \text{deltaf} \cdot \frac{1}{1 + \exp(1 \cdot (sr) \cdot (time - to + t1 - 3))} - stim \cdot CBF0_Fin_t \cdot \text{deltaf} \cdot \frac{1}{1 + \exp(1 \cdot (sr) \cdot (time - (to + tend + t1 + 3)))} \right) \quad (50)$$

6.51 Rule V_{c_O2}

Rule V_{c_O2} is an assignment rule for parameter V_{c_O2} :

$$V_{c_O2} = 2 \cdot \frac{Fin_t}{V_c} \cdot ([O2a] - [O2c]) \quad (51)$$

6.52 Rule Vc_GLC

Rule Vc_GLC is an assignment rule for parameter Vc_GLC :

$$Vc_GLC = 2 \cdot \frac{Fin_t}{Vc} \cdot ([GLCa] - [GLCc]) \quad (52)$$

6.53 Rule Vc_LAC

Rule Vc_LAC is an assignment rule for parameter Vc_LAC :

$$Vc_LAC = 2 \cdot \frac{Fin_t}{Vc} \cdot ([LACa] - [LACc]) \quad (53)$$

6.54 Rule Vc_CO2

Rule Vc_CO2 is an assignment rule for parameter Vc_CO2 :

$$Vc_CO2 = 2 \cdot \frac{Fin_t}{Vc} \cdot ([CO2c] - [CO2a]) \quad (54)$$

6.55 Rule $Fout_t$

Rule $Fout_t$ is an assignment rule for parameter $Fout_t$:

$$Fout_t = CBF0_model_parameters \cdot \frac{\left(\frac{Vv}{Vv0}\right)^2 + tv \cdot \left(\frac{Vv}{Vv0}\right)^{0.5} \cdot \frac{Fin_t}{Vv0}}{1 + CBF0_model_parameters \cdot tv \cdot \left(\frac{Vv}{Vv0}\right)^{0.5} \cdot \frac{1}{Vv0}} \quad (55)$$

6.56 Rule $BOLD$

Rule $BOLD$ is an assignment rule for parameter $BOLD$:

$$BOLD = Vv0 \cdot \left((k1 + k2) \cdot \left(1 - \frac{dHb}{dHb0} \right) - (k2 + k3) \cdot \left(1 - \frac{Vv}{Vv0} \right) \right) \quad (56)$$

6.57 Rule $unitpulseSB$

Rule $unitpulseSB$ is an assignment rule for parameter $unitpulseSB$:

$$unitpulseSB = \begin{cases} 1 & \text{if } (time \geq to) \wedge (time \leq to + tend) \\ 0 & \text{otherwise} \end{cases} \quad (57)$$

6.58 Rule v_stim

Rule v_stim is an assignment rule for parameter v_stim :

$$v_stim = stim \cdot \left(v1_n + v2_n \cdot \frac{time - to}{t_n_stim} \cdot \exp \left(\left((time - to) \cdot \frac{unitpulseSB}{t_n_stim} \right) \right) \right) \cdot unitpulseSB \quad (58)$$

6.59 Rule Vn_stim

Rule Vn_stim is an assignment rule for parameter Vn_stim :

$$Vn_stim = v_stim \quad (59)$$

6.60 Rule Vn_stim_GLU

Rule Vn_stim_GLU is an assignment rule for parameter Vn_stim_GLU :

$$Vn_stim_GLU = Vn_stim \cdot R_GLU_NA \cdot KO2 \cdot \frac{[GLUn]}{[GLUn] + Km_GLU} \quad (60)$$

6.61 Rule $unitstepSB$

Rule $unitstepSB$ is an assignment rule for parameter $unitstepSB$:

$$unitstepSB = \begin{cases} 1 & \text{if } time - (tend + to) \geq 0 \\ 0 & \text{otherwise} \end{cases} \quad (61)$$

6.62 Rule $unitstepSB2$

Rule $unitstepSB2$ is an assignment rule for parameter $unitstepSB2$:

$$unitstepSB2 = \begin{cases} 1 & \text{if } time - (tend_GLY + to + to_GLY) \geq 0 \\ 0 & \text{otherwise} \end{cases} \quad (62)$$

6.63 Rule $deltaVt_GLY$

Rule $deltaVt_GLY$ is an assignment rule for parameter $deltaVt_GLY$:

$$\begin{aligned} deltaVt_GLY = & 1 + stim \cdot delta_GLY \cdot KO3 \\ & \cdot \frac{1}{1 + \exp(1 \cdot (sr_GLY) \cdot (time - (to + to_GLY)))} \cdot (1 - unitstepSB2) \end{aligned} \quad (63)$$

6.64 Rule Vg_glyp

Rule Vg_glyp is an assignment rule for parameter Vg_glyp :

$$Vg_glyp = Vmax_glyp \cdot \frac{[GLYg]}{[GLYg] + Km_GLY} \cdot deltaVt_GLY \quad (64)$$

6.65 Rule NAg

Rule NAg is a rate rule for species NAg :

$$\frac{d}{dt}NAg = Vg_leak_Na + 3 \cdot Veg_GLU - 3 \cdot Vg_pump \quad (65)$$

6.66 Rule GLCg

Rule GLCg is a rate rule for species GLCg:

$$\frac{d}{dt}\text{GLCg} = V_{\text{cg_GLC}} + V_{\text{eg_GLC}} - V_{\text{g_hk}} \quad (66)$$

6.67 Rule G6Pg

Rule G6Pg is a rate rule for species G6Pg:

$$\frac{d}{dt}\text{G6Pg} = V_{\text{g_hk}} + V_{\text{g_glyp}} - (V_{\text{g_pgi}} + V_{\text{g_glys}}) \quad (67)$$

6.68 Rule F6Pg

Rule F6Pg is a rate rule for species F6Pg:

$$\frac{d}{dt}\text{F6Pg} = V_{\text{g_pgi}} - V_{\text{g_pfk}} \quad (68)$$

6.69 Rule GAPg

Rule GAPg is a rate rule for species GAPg:

$$\frac{d}{dt}\text{GAPg} = 2 \cdot V_{\text{g_pfk}} - V_{\text{g_pgk}} \quad (69)$$

6.70 Rule PEPg

Rule PEPg is a rate rule for species PEPg:

$$\frac{d}{dt}\text{PEPg} = V_{\text{g_pgk}} - V_{\text{g_pk}} \quad (70)$$

6.71 Rule PYRg

Rule PYRg is a rate rule for species PYRg:

$$\frac{d}{dt}\text{PYRg} = V_{\text{g_pk}} - (V_{\text{g_ldh}} + V_{\text{g_mito}}) \quad (71)$$

6.72 Rule LACg

Rule LACg is a rate rule for species LACg:

$$\frac{d}{dt}\text{LACg} = V_{\text{g_ldh}} - (V_{\text{ge_LAC}} + V_{\text{gc_LAC}}) \quad (72)$$

6.73 Rule NADHg

Rule NADHg is a rate rule for species NADHg:

$$\frac{d}{dt}\text{NADHg} = V_{g_pgk} - (V_{g_ldh} + V_{g_mito}) \quad (73)$$

6.74 Rule ATPg

Rule ATPg is a rate rule for species ATPg:

$$\begin{aligned} \frac{d}{dt}\text{ATPg} = & (V_{g_pgk} + V_{g_pk} + nOP \cdot V_{g_mito} + V_{g_ck} \\ & - (V_{g_hk} + V_{g_pfk} + V_{g_ATPase} + V_{g_pump} + V_{g_gs})) \cdot (1 - dAMP_dATPg)^1 \end{aligned} \quad (74)$$

6.75 Rule PCrg

Rule PCrg is a rate rule for species PCrg:

$$\frac{d}{dt}\text{PCrg} = V_{g_ck} \quad (75)$$

6.76 Rule O2g

Rule O2g is a rate rule for species O2g:

$$\frac{d}{dt}\text{O2g} = V_{cg_O2} - NAero \cdot V_{g_mito} \quad (76)$$

6.77 Rule GLYg

Rule GLYg is a rate rule for species GLYg:

$$\frac{d}{dt}\text{GLYg} = V_{g_glys} - V_{g_glyp} \quad (77)$$

6.78 Rule GLUg

Rule GLUg is a rate rule for species GLUg:

$$\frac{d}{dt}\text{GLUg} = V_{eg_GLU} - V_{g_gs} \quad (78)$$

6.79 Rule O2c

Rule O2c is a rate rule for species O2c:

$$\frac{d}{dt}\text{O2c} = V_{c_O2} - \left(V_{cn_O2} \cdot \frac{1}{R_{cn}} + V_{cg_O2} \cdot \frac{1}{R_{cg}} \right) \quad (79)$$

6.80 Rule GLCc

Rule GLCc is a rate rule for species GLCc:

$$\frac{d}{dt}\text{GLCc} = V_{c_GLC} - \left(V_{ce_GLC} \cdot \frac{1}{R_{ce}} + V_{cg_GLC} \cdot \frac{1}{R_{cg}} \right) \quad (80)$$

6.81 Rule LACc

Rule LACc is a rate rule for species LACc:

$$\frac{d}{dt}\text{LACc} = V_{c_LAC} + V_{ec_LAC} \cdot \frac{1}{R_{ce}} + V_{gc_LAC} \cdot \frac{1}{R_{cg}} \quad (81)$$

6.82 Rule CO2c

Rule CO2c is a rate rule for species CO2c:

$$\frac{d}{dt}\text{CO2c} = V_{nc_CO2} \cdot \frac{1}{R_{cn}} + V_{gc_CO2} \cdot \frac{1}{R_{cg}} - V_{c_CO2} \quad (82)$$

6.83 Rule GLCe

Rule GLCe is a rate rule for species GLCe:

$$\frac{d}{dt}\text{GLCe} = V_{ce_GLC} - \left(V_{eg_GLC} \cdot \frac{1}{R_{eg}} + V_{en_GLC} \cdot \frac{1}{R_{en}} \right) \quad (83)$$

6.84 Rule LACe

Rule LACe is a rate rule for species LACe:

$$\frac{d}{dt}\text{LACe} = V_{ne_LAC} \cdot \frac{1}{R_{en}} + V_{ge_LAC} \cdot \frac{1}{R_{eg}} - V_{ec_LAC} \quad (84)$$

6.85 Rule GLUe

Rule GLUe is a rate rule for species GLUe:

$$\frac{d}{dt}\text{GLUe} = V_{n_stim_GLU} \cdot \frac{1}{R_{en}} - V_{eg_GLU} \cdot \frac{1}{R_{eg}} \quad (85)$$

6.86 Rule NAn

Rule NAn is a rate rule for species NAn:

$$\frac{d}{dt}\text{NAn} = V_{n_leak_Na} + V_{n_stim} - 3 \cdot V_{n_pump} \quad (86)$$

6.87 Rule GLCn

Rule GLCn is a rate rule for species GLCn:

$$\frac{d}{dt} \text{GLCn} = V_{\text{en_GLC}} - V_{\text{n_hk}} \quad (87)$$

6.88 Rule G6Pn

Rule G6Pn is a rate rule for species G6Pn:

$$\frac{d}{dt} \text{G6Pn} = V_{\text{n_hk}} - V_{\text{n_pgi}} \quad (88)$$

6.89 Rule F6Pn

Rule F6Pn is a rate rule for species F6Pn:

$$\frac{d}{dt} \text{F6Pn} = V_{\text{n_pgi}} - V_{\text{n_pfk}} \quad (89)$$

6.90 Rule GAPn

Rule GAPn is a rate rule for species GAPn:

$$\frac{d}{dt} \text{GAPn} = 2 \cdot V_{\text{n_pfk}} - V_{\text{n_pgk}} \quad (90)$$

6.91 Rule PEPn

Rule PEPn is a rate rule for species PEPn:

$$\frac{d}{dt} \text{PEPn} = V_{\text{n_pgk}} - V_{\text{n_pk}} \quad (91)$$

6.92 Rule PYRn

Rule PYRn is a rate rule for species PYRn:

$$\frac{d}{dt} \text{PYRn} = V_{\text{n_pk}} - (V_{\text{n_ldh}} + V_{\text{n_mito}}) \quad (92)$$

6.93 Rule LACn

Rule LACn is a rate rule for species LACn:

$$\frac{d}{dt} \text{LACn} = V_{\text{n_ldh}} - V_{\text{ne_LAC}} \quad (93)$$

6.94 Rule NADHn

Rule NADHn is a rate rule for species NADHn:

$$\frac{d}{dt}\text{NADHn} = \text{Vn_pgk} - (\text{Vn_ldh} + \text{Vn_mito}) \quad (94)$$

6.95 Rule ATPn

Rule ATPn is a rate rule for species ATPn:

$$\begin{aligned} \frac{d}{dt}\text{ATPn} = & (\text{Vn_pgk} + \text{Vn_pk} + \text{nOP} \cdot \text{Vn_mito} + \text{Vn_ck} \\ & - (\text{Vn_hk} + \text{Vn_pfk} + \text{Vn_ATPase} + \text{Vn_pump})) \cdot (1 - \text{dAMP_dATPn})^1 \end{aligned} \quad (95)$$

6.96 Rule PCrn

Rule PCrn is a rate rule for species PCrn:

$$\frac{d}{dt}\text{PCrn} = \text{Vn_ck} \quad (96)$$

6.97 Rule O2n

Rule O2n is a rate rule for species O2n:

$$\frac{d}{dt}\text{O2n} = \text{Vcn_O2} - \text{NAero} \cdot \text{Vn_mito} \quad (97)$$

6.98 Rule GLUn

Rule GLUn is a rate rule for species GLUn:

$$\frac{d}{dt}\text{GLUn} = \text{Vg_gs} \cdot \frac{1}{\text{Rng}} - \text{Vn_stim_GLU} \quad (98)$$

6.99 Rule Vv

Rule Vv is a rate rule for parameter Vv:

$$\frac{d}{dt}\text{Vv} = \text{Fin_t} - \text{Fout_t} \quad (99)$$

6.100 Rule dHb

Rule dHb is a rate rule for parameter dHb:

$$\frac{d}{dt}\text{dHb} = \text{Fin_t} \cdot ([\text{O2a}] - [\text{O2c}]) - \text{Fout_t} \cdot \frac{\text{dHb}}{\text{Vv}} \quad (100)$$

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 [N_{Ag}](#)

Name N_{Ag}

Notes Sodium

Initial concentration 13.36 mol · l⁻¹

Involved in rule [N_{Ag}](#)

One rule determines the species' quantity.

7.2 Species [GLC_g](#)

Name GLC_g

Notes Glucose

Initial concentration 0.1656 mol · l⁻¹

Involved in rule [GLC_g](#)

One rule determines the species' quantity.

7.3 Species [G6P_g](#)

Name G6P_g

Notes Glucose-6-P

Initial concentration 0.7326 mol · l⁻¹

Involved in rule [G6P_g](#)

One rule determines the species' quantity.

7.4 Species [F6P_g](#)

Name F6P_g

Notes Fructose-6-P

Initial concentration 0.1116 mol · l⁻¹

Involved in rule [F6P_g](#)

One rule determines the species' quantity.

7.5 Species GAPg

Name GAPg

Notes Glyceraldehyde-3-P

Initial concentration $0.0698 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule GAPg

One rule determines the species' quantity.

7.6 Species PEPg

Name PEPg

Notes Phosphoenolpyruvate

Initial concentration $0.0254 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule PEPg

One rule determines the species' quantity.

7.7 Species PYRg

Name PYRg

Notes Pyruvate

Initial concentration $0.1711 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule PYRg

One rule determines the species' quantity.

7.8 Species LACg

Name LACg

Notes Lactate

Initial concentration $0.4651 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule LACg

One rule determines the species' quantity.

7.9 Species NADHg

Name NADHg

Notes Nicotinamide adenine dinucleotide reduced

Initial concentration $0.0445 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule NADHg

One rule determines the species' quantity.

7.10 Species ATPg

Name ATPg

Notes Adenosine triphosphate

Initial concentration $2.24 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule ATPg

One rule determines the species' quantity.

7.11 Species PCrg

Name PCrg

Notes Phosphocreatine

Initial concentration $4.6817 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule PCrg

One rule determines the species' quantity.

7.12 Species O2g

Name O2g

Notes Oxygen

Initial concentration $0.1589 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule O2g

One rule determines the species' quantity.

7.13 Species GLYg

Name GLYg

Notes Glycogen

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

Involved in rule GLYg

One rule determines the species' quantity.

7.14 Species GLUg

Name GLUg

Notes Glutamate

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

Involved in rule GLUg

One rule determines the species' quantity.

7.15 Species NADg

Name NADg

Notes Nicotinamide adenine dinucleotide oxidized

Initial concentration $0.1755 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule NADg

One rule determines the species' quantity.

7.16 Species ADPg

Name ADPg

Notes Adenosine diphosphate

Initial concentration $0.13070953832961 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule ADPg

One rule determines the species' quantity.

7.17 Species CR_g

Name CR_g

Notes Creatine

Initial concentration 0.318300000000001 mol · l⁻¹

Involved in rule CR_g

One rule determines the species' quantity.

7.18 Species NADH_{g-tot}

Name NADH_{g-tot}

Notes Nicotinamide adenine dinucleotide reduced

Initial concentration 0.22 mol · l⁻¹

$$\frac{d}{dt} \text{NADH}_{g\text{-tot}} = 0 \quad (101)$$

7.19 Species PCr_{g-tot}

Name PCr_{g-tot}

Notes Phosphocreatine

Initial concentration 5 mol · l⁻¹

$$\frac{d}{dt} \text{PCr}_{g\text{-tot}} = 0 \quad (102)$$

7.20 Species AMP_g

Name AMP_g

Notes Adenosine monophosphate

Initial concentration 0.00829046167039005 mol · l⁻¹

Involved in rule AMP_g

One rule determines the species' quantity.

7.21 Species [O2c](#)

Name O2c

Notes Oxygen

Initial concentration $7.4201 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule [O2c](#)

One rule determines the species' quantity.

7.22 Species [GLCc](#)

Name GLCc

Notes Glucose

Initial concentration $4.6401 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule [GLCc](#)

One rule determines the species' quantity.

7.23 Species [LACc](#)

Name LACc

Notes Lactate

Initial concentration $0.3251 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule [LACc](#)

One rule determines the species' quantity.

7.24 Species [CO2c](#)

Name CO2c

Notes Carbon dioxide

Initial concentration $2.12 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule [CO2c](#)

One rule determines the species' quantity.

7.25 Species GLCe

Name GLCe

Notes Glucose

Initial concentration $0.3339 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule GLCe

One rule determines the species' quantity.

7.26 Species LACe

Name LACe

Notes Lactate

Initial concentration $0.3986 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule LACe

One rule determines the species' quantity.

7.27 Species GLUe

Name GLUe

Notes Glutamate

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

Involved in rule GLUe

One rule determines the species' quantity.

7.28 Species NAe

Name NAe

Notes Sodium

Initial concentration $150 \text{ mol} \cdot \text{l}^{-1}$

$$\frac{d}{dt} \text{NAe} = 0 \quad (103)$$

7.29 Species O2a

Name O2a

Notes Oxygen

Initial concentration $8.34 \text{ mol} \cdot \text{l}^{-1}$

$$\frac{d}{dt} \text{O2a} = 0 \quad (104)$$

7.30 Species CO2a

Name CO2a

Notes Carbon dioxide

Initial concentration $1.2 \text{ mol} \cdot \text{l}^{-1}$

$$\frac{d}{dt} \text{CO2a} = 0 \quad (105)$$

7.31 Species GLCa

Name GLCa

Notes Glucose

Initial concentration $4.8 \text{ mol} \cdot \text{l}^{-1}$

$$\frac{d}{dt} \text{GLCa} = 0 \quad (106)$$

7.32 Species LACa

Name LACa

Notes Lactate

Initial concentration $0.313 \text{ mol} \cdot \text{l}^{-1}$

$$\frac{d}{dt} \text{LACa} = 0 \quad (107)$$

7.33 Species [NAn](#)

Name NAn

Notes Sodium

Initial concentration $15.533 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule [NAn](#)

One rule determines the species' quantity.

7.34 Species [GLCn](#)

Name GLCn

Notes Glucose

Initial concentration $0.2633 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule [GLCn](#)

One rule determines the species' quantity.

7.35 Species [G6Pn](#)

Name G6Pn

Notes Glucose-6-P

Initial concentration $0.7275 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule [G6Pn](#)

One rule determines the species' quantity.

7.36 Species [F6Pn](#)

Name F6Pn

Notes Fructose-6-P

Initial concentration $0.1091 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule [F6Pn](#)

One rule determines the species' quantity.

7.37 Species [GAPn](#)

Name GAPn

Notes Glyceraldehyde-3-P

Initial concentration $0.0418 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule [GAPn](#)

One rule determines the species' quantity.

7.38 Species [PEPn](#)

Name PEPn

Notes Phosphoenolpyruvate

Initial concentration $0.0037 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule [PEPn](#)

One rule determines the species' quantity.

7.39 Species [PYRn](#)

Name PYRn

Notes Pyruvate

Initial concentration $0.0388 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule [PYRn](#)

One rule determines the species' quantity.

7.40 Species [LACn](#)

Name LACn

Notes Lactate

Initial concentration $0.3856 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule [LACn](#)

One rule determines the species' quantity.

7.41 Species [NADHn](#)

Name NADHn

Notes Nicotinamide adenine dinucleotide reduced

Initial concentration $0.0319 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule [NADHn](#)

One rule determines the species' quantity.

7.42 Species [ATPn](#)

Name ATPn

Notes Adenosine triphosphate

Initial concentration $2.2592 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule [ATPn](#)

One rule determines the species' quantity.

7.43 Species [PCrn](#)

Name PCrn

Notes Posphocreatine

Initial concentration $4.2529 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule [PCrn](#)

One rule determines the species' quantity.

7.44 Species [O2n](#)

Name O2n

Notes Oxygen

Initial concentration $0.0975 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule [O2n](#)

One rule determines the species' quantity.

7.45 Species [GLUn](#)

Name GLUn

Notes Glutamate

Initial concentration $3 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule [GLUn](#)

One rule determines the species' quantity.

7.46 Species [ADPn](#)

Name ADPn

Notes Adenosine diphosphate

Initial concentration $0.113591983539553 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule [ADPn](#)

One rule determines the species' quantity.

7.47 Species [CRn](#)

Name CRn

Notes Creatine

Initial concentration $0.7471 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule [CRn](#)

One rule determines the species' quantity.

7.48 Species [NADn](#)

Name NADn

Notes Nicotinamide adenine dinucleotide oxidized

Initial concentration $0.1881 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule [NADn](#)

One rule determines the species' quantity.

7.49 Species `NADH_n_tot`

Name `NADH_n_tot`

Notes Nicotinamide adenine dinucleotide reduced

Initial concentration $0.22 \text{ mol} \cdot \text{l}^{-1}$

$$\frac{d}{dt} \text{NADH_n_tot} = 0 \quad (108)$$

7.50 Species `PCrn_tot`

Name `PCrn_tot`

Notes Phosphocreatine

Initial concentration $5 \text{ mol} \cdot \text{l}^{-1}$

$$\frac{d}{dt} \text{PCrn_tot} = 0 \quad (109)$$

7.51 Species `AMPn`

Name `AMPn`

Notes Adenosine monophosphate

Initial concentration $0.006208016460449 \text{ mol} \cdot \text{l}^{-1}$

Involved in rule `AMPn`

One rule determines the species' quantity.

SBML2^{LaTeX} 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