

SBML Model Report

Model name: “Noble1998_VentricularCellModel_ModelC”



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: Penny Noble¹, Penny Noble² and Penny Noble³ at February second 2012 at 1:30 p. m. and last time modified at February second 2012 at 9:19 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	229	unit definitions	18
rules	106	initial assignments	0

Model Notes

This a model from the article:

Improved guinea-pig ventricular cell model incorporating a diadic space, IKr andIKs, and length- and tension-dependent processes.

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Noble D, Varghese A, Kohl P, Noble P. [Can J Cardiol](#) 1998 Jan;14(1):123-34 [9487284](#) ,

Abstract:

The guinea-pig ventricular cell model, originally developed by Noble et al in 1991, has been greatly extended to include accumulation and depletion of calcium in a diadic space between the sarcolemma and the sarcoplasmic reticulum where, according to contemporary understanding, the majority of calcium-induced calcium release is triggered. The calcium in this space is also assumed to play the major role in calcium-induced inactivation of the calcium current. Delayed potassium current equations have been developed to include the rapid (IKr) and slow (IKs) components of the delayed rectifier current based on the data of Heath and Terrar, along with data from Sanguinetti and Jurkiewicz. Length- and tension-dependent changes in mechanical and electrophysiological processes have been incorporated as described recently by Kohl et al. Drug receptor interactions have started to be developed, using the sodium channel as the first target. The new model has been tested against experimental data on action potential clamp, and on force-interval and duration-interval relations; it has been found to reliably reproduce experimental observations.

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

The original model was: [Noble D, Varghese A, Kohl P, Noble P. \(1998\) - 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 23 unit definitions of which five are predefined by SBML and not mentioned in the model.

2.1 Unit micrometre

Name micrometre

Definition μm

2.2 Unit per_micrometre

Name per_micrometre

Definition μm^{-1}

2.3 Unit per_second

Name per_second

Definition s^{-1}

2.4 Unit millivolt

Name millivolt

Definition mV

2.5 Unit per_millivolt

Name per_millivolt

Definition mV^{-1}

2.6 Unit per_millivolt_second

Name per_millivolt_second

Definition $\text{mV}^{-1} \cdot \text{s}^{-1}$

2.7 Unit microS

Name microS

Definition μS

2.8 Unit microF

Name microF

Definition μF

2.9 Unit nanoA

Name nanoA

Definition nA

2.10 Unit nanoA_per_millimolar

Name nanoA_per_millimolar

Definition nA · mmol⁻¹ · l

2.11 Unit millimolar

Name millimolar

Definition mmol · l⁻¹

2.12 Unit millimolar4

Name millimolar4

Definition mmol⁴ · l⁻⁴

2.13 Unit millimolar_per_second

Name millimolar_per_second

Definition mmol · l⁻¹ · s⁻¹

2.14 Unit per_millimolar_second

Name per_millimolar_second

Definition mmol⁻¹ · l · s⁻¹

2.15 Unit joule_per_kilomole_kelvin

Name joule_per_kilomole_kelvin

Definition J · kmol⁻¹ · K⁻¹

2.16 Unit coulomb_per_mole

Name coulomb_per_mole

Definition C · mol⁻¹

2.17 Unit `micrometre3`

Name `micrometre3`

Definition μm^3

2.18 Unit `litre_micrometre3`

Name `litre_micrometre3`

Definition $1 \cdot \mu\text{m}^{-3}$

2.19 Unit `substance`

Notes Mole is the predefined SBML unit for `substance`.

Definition `mol`

2.20 Unit `volume`

Notes Litre is the predefined SBML unit for `volume`.

Definition `l`

2.21 Unit `area`

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

Definition m^2

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 Compartment

This model contains one compartment.

Table 2: Properties of all compartments.

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

3.1 Compartment COMpartment

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

4 Parameters

This model contains 229 global parameters.

Table 3: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
V_membrane	V		-92.849		<input type="checkbox"/>
R	R		8314.472		<input checked="" type="checkbox"/>
T	T		310.000		<input checked="" type="checkbox"/>
F	F		96485.342		<input checked="" type="checkbox"/>
Cm	Cm		$9.5 \cdot 10^{-5}$		<input checked="" type="checkbox"/>
i_Stim	i_Stim		0.000		<input type="checkbox"/>
stim_start	stim_start		0.100		<input checked="" type="checkbox"/>
stim_end	stim_end		10000.000		<input checked="" type="checkbox"/>
stim_period	stim_period		1.000		<input checked="" type="checkbox"/>
stim- _duration	stim_duration		0.003		<input checked="" type="checkbox"/>
stim- _amplitude	stim_amplitude		-3.000		<input checked="" type="checkbox"/>
E_Na	E_Na		0.000		<input type="checkbox"/>
E_K	E_K		0.000		<input type="checkbox"/>
E_Ks	E_Ks		0.000		<input type="checkbox"/>
E_Ca	E_Ca		0.000		<input type="checkbox"/>
E_mh	E_mh		0.000		<input type="checkbox"/>
P_kna	P_kna		0.030		<input checked="" type="checkbox"/>
i_K1	i_K1		0.000		<input type="checkbox"/>
K_mk1	K_mk1		10.000		<input checked="" type="checkbox"/>
g_K1	g_K1		0.500		<input checked="" type="checkbox"/>
i_Kr	i_Kr		0.000		<input type="checkbox"/>
g_Kr1	g_Kr1		0.002		<input checked="" type="checkbox"/>
g_Kr2	g_Kr2		0.001		<input checked="" type="checkbox"/>

Id	Name	SBO	Value	Unit	Constant
xr1	xr1		$1.03 \cdot 10^{-5}$		<input type="checkbox"/>
V_rapid- _delayed- _rectifier- _potassium- _current_xr1- _gate	V		0.000		<input checked="" type="checkbox"/>
alpha_xr1	alpha_xr1		0.000		<input type="checkbox"/>
beta_xr1	beta_xr1		0.000		<input type="checkbox"/>
time_rapid- _delayed- _rectifier- _potassium- _current_xr1- _gate	time		0.000		<input checked="" type="checkbox"/>
xr2	xr2		$2 \cdot 10^{-7}$		<input type="checkbox"/>
V_rapid- _delayed- _rectifier- _potassium- _current_xr2- _gate	V		0.000		<input checked="" type="checkbox"/>
alpha_xr2	alpha_xr2		0.000		<input type="checkbox"/>
beta_xr2	beta_xr2		0.000		<input type="checkbox"/>
time_rapid- _delayed- _rectifier- _potassium- _current_xr2- _gate	time		0.000		<input checked="" type="checkbox"/>
i_Ks	i_Ks		0.000		<input type="checkbox"/>
g_Ks	g_Ks		0.003		<input checked="" type="checkbox"/>
xs	xs		0.001		<input type="checkbox"/>
V_slow- _delayed- _rectifier- _potassium- _current_xs- _gate	V		0.000		<input checked="" type="checkbox"/>
alpha_xs	alpha_xs		0.000		<input type="checkbox"/>
beta_xs	beta_xs		0.000		<input type="checkbox"/>

Id	Name	SBO	Value	Unit	Constant
time_slow- _delayed- _rectifier- _potassium- _current_xs- _gate	time		0.000		<input checked="" type="checkbox"/>
i_K_ATP	i_K_ATP		0.000		<input type="checkbox"/>
g_K_ATP	g_K_ATP		0.000		<input checked="" type="checkbox"/>
K_ATP	K_ATP		0.100		<input checked="" type="checkbox"/>
ATP	ATP		0.000		<input checked="" type="checkbox"/>
i_KNa	i_KNa		0.000		<input type="checkbox"/>
g_K_Na	g_K_Na		0.000		<input checked="" type="checkbox"/>
K_kna	K_kna		20.000		<input checked="" type="checkbox"/>
i_Na	i_Na		0.000		<input type="checkbox"/>
g_Na	g_Na		2.500		<input checked="" type="checkbox"/>
m	m		0.002		<input type="checkbox"/>
alpha_m	alpha_m		0.000		<input type="checkbox"/>
beta_m	beta_m		0.000		<input type="checkbox"/>
V_fast-	V		0.000		<input checked="" type="checkbox"/>
_sodium- _current_m- _gate					
time_fast- _sodium- _current_m- _gate	time		0.000		<input checked="" type="checkbox"/>
delta_m	delta_m		10^{-5}		<input checked="" type="checkbox"/>
E0_m	E0_m		0.000		<input type="checkbox"/>
h	h		0.994		<input type="checkbox"/>
alpha_h	alpha_h		0.000		<input type="checkbox"/>
beta_h	beta_h		0.000		<input type="checkbox"/>
V_fast-	V		0.000		<input checked="" type="checkbox"/>
_sodium- _current_h- _gate					
time_fast- _sodium- _current_h- _gate	time		0.000		<input checked="" type="checkbox"/>
i_p_Na	i_p_Na		0.000		<input type="checkbox"/>
g_pna	g_pna		0.004		<input checked="" type="checkbox"/>
i_b_Na	i_b_Na		0.000		<input type="checkbox"/>

Id	Name	SBO	Value	Unit	Constant
g_bna	g_bna		$6 \cdot 10^{-4}$		<input checked="" type="checkbox"/>
i_Ca_L_Ca_cyt	i_Ca_L_Ca_cyt		0.000		<input type="checkbox"/>
i_Ca_L_K_cyt	i_Ca_L_K_cyt		0.000		<input type="checkbox"/>
i_Ca_L_Na_cyt	i_Ca_L_Na_cyt		0.000		<input type="checkbox"/>
i_Ca_L_Ca_ds	i_Ca_L_Ca_ds		0.000		<input type="checkbox"/>
i_Ca_L_K_ds	i_Ca_L_K_ds		0.000		<input type="checkbox"/>
i_Ca_L_Na_ds	i_Ca_L_Na_ds		0.000		<input type="checkbox"/>
i_Ca_L	i_Ca_L		0.000		<input type="checkbox"/>
P_Ca_L	P_Ca_L		0.100		<input checked="" type="checkbox"/>
P_Ca_K	P_Ca_K		0.002		<input checked="" type="checkbox"/>
P_Ca_Na	P_Ca_Na		0.010		<input checked="" type="checkbox"/>
Km_f2	Km_f2		100000.000		<input checked="" type="checkbox"/>
Km_f2ds	Km_f2ds		0.001		<input checked="" type="checkbox"/>
R_decay	R_decay		20.000		<input checked="" type="checkbox"/>
FrICa	FrICa		1.000		<input checked="" type="checkbox"/>
d	d		0.000		<input type="checkbox"/>
alpha_d	alpha_d		0.000		<input type="checkbox"/>
beta_d	beta_d		0.000		<input type="checkbox"/>
V_L.type_Ca- _channel_d- _gate	V		0.000		<input checked="" type="checkbox"/>
time_L.type- _Ca_channel- _d_gate	time		0.000		<input checked="" type="checkbox"/>
E0_d	E0_d		0.000		<input type="checkbox"/>
speed_d	speed_d		3.000		<input checked="" type="checkbox"/>
f	f		1.000		<input type="checkbox"/>
alpha_f	alpha_f		0.000		<input type="checkbox"/>
beta_f	beta_f		0.000		<input type="checkbox"/>
V_L.type_Ca- _channel_f- _gate	V		0.000		<input checked="" type="checkbox"/>
time_L.type- _Ca_channel- _f_gate	time		0.000		<input checked="" type="checkbox"/>
speed_f	speed_f		0.300		<input checked="" type="checkbox"/>
delta_f	delta_f		10^{-4}		<input checked="" type="checkbox"/>
E0_f	E0_f		0.000		<input type="checkbox"/>
f2	f2		0.935		<input type="checkbox"/>
Ca.i.L.type- _Ca_channel- _f2_gate	Ca.i		0.000		<input checked="" type="checkbox"/>

Id	Name	SBO	Value	Unit	Constant
V.L.type-Ca- _channel_f2- _gate	V		0.000		<input checked="" type="checkbox"/>
time.L.type- _Ca_channel- _f2_gate	time		0.000		<input checked="" type="checkbox"/>
f2ds	f2ds		0.965		<input type="checkbox"/>
Ca_ds.L.type- _Ca_channel- _f2ds_gate	Ca_ds		0.000		<input checked="" type="checkbox"/>
V.L.type-Ca- _channel- _f2ds_gate	V		0.000		<input checked="" type="checkbox"/>
time.L.type- _Ca_channel- _f2ds_gate	time		0.000		<input checked="" type="checkbox"/>
i_b.Ca	i_b.Ca		0.000		<input type="checkbox"/>
g_bca	g_bca		$2.5 \cdot 10^{-4}$		<input checked="" type="checkbox"/>
i_to	i_to		0.000		<input type="checkbox"/>
g_to	g_to		0.005		<input checked="" type="checkbox"/>
g_tos	g_tos		0.000		<input checked="" type="checkbox"/>
s	s		0.995		<input type="checkbox"/>
alpha_s	alpha_s		0.000		<input type="checkbox"/>
beta_s	beta_s		0.000		<input type="checkbox"/>
V.transient- _outward- _current_s- _gate	V		0.000		<input checked="" type="checkbox"/>
time- _transient- _outward- _current_s- _gate	time		0.000		<input checked="" type="checkbox"/>
r	r		0.000		<input type="checkbox"/>
V.transient- _outward- _current_r- _gate	V		0.000		<input checked="" type="checkbox"/>

Id	Name	SBO	Value	Unit	Constant
time- _transient- _outward- _current_r- _gate	time		0.000		<input checked="" type="checkbox"/>
i_K_ACh	i_K_ACh		0.000		<input type="checkbox"/>
g_KACh	g_KACh		0.000		<input checked="" type="checkbox"/>
ACh	ACh		5.000		<input checked="" type="checkbox"/>
K_D	K_D		$1.3 \cdot 10^{-4}$		<input checked="" type="checkbox"/>
x_ACh	x_ACh		0.000		<input type="checkbox"/>
alpha_ACh	alpha_ACh		0.500		<input checked="" type="checkbox"/>
beta_ACh	beta_ACh		0.500		<input checked="" type="checkbox"/>
time_ACh- _dependent- _potassium- _current- _x_ACh_gate	time		0.000		<input checked="" type="checkbox"/>
i_NaK	i_NaK		0.000		<input type="checkbox"/>
i_NaK_max	i_NaK_max		0.700		<input checked="" type="checkbox"/>
K_mK	K_mK		1.000		<input checked="" type="checkbox"/>
K_mNa	K_mNa		40.000		<input checked="" type="checkbox"/>
i_NaCa	i_NaCa		0.000		<input type="checkbox"/>
i_NaCa_cyt	i_NaCa_cyt		0.000		<input type="checkbox"/>
i_NaCa_ds	i_NaCa_ds		0.000		<input type="checkbox"/>
k_NaCa	k_NaCa		$5 \cdot 10^{-4}$		<input checked="" type="checkbox"/>
n_NaCa	n_NaCa		3.000		<input checked="" type="checkbox"/>
d_NaCa	d_NaCa		0.000		<input checked="" type="checkbox"/>
gamma	gamma		0.500		<input checked="" type="checkbox"/>
FRiNaCa	FRiNaCa		0.001		<input checked="" type="checkbox"/>
i_up	i_up		0.000		<input type="checkbox"/>
K_1	K_1		0.000		<input type="checkbox"/>
K_2	K_2		0.000		<input type="checkbox"/>
K_cyca	K_cyca		$3 \cdot 10^{-4}$		<input checked="" type="checkbox"/>
K_xcs	K_xcs		0.400		<input checked="" type="checkbox"/>
K_srca	K_srca		0.500		<input checked="" type="checkbox"/>
alpha_up	alpha_up		0.400		<input checked="" type="checkbox"/>
beta_up	beta_up		0.030		<input checked="" type="checkbox"/>
i_trans	i_trans		0.000		<input type="checkbox"/>
i_rel	i_rel		0.000		<input type="checkbox"/>
VoltDep	VoltDep		0.000		<input type="checkbox"/>
RegBindSite	RegBindSite		0.000		<input type="checkbox"/>
CaiReg	CaiReg		0.000		<input type="checkbox"/>

Id	Name	SBO	Value	Unit	Constant
CadsReg	CadsReg		0.000		<input type="checkbox"/>
ActRate	ActRate		0.000		<input type="checkbox"/>
InactRate	InactRate		0.000		<input type="checkbox"/>
K_leak_rate	K_leak_rate		0.005		<input checked="" type="checkbox"/>
SRLeak	SRLeak		0.000		<input type="checkbox"/>
K_m_rel	K_m_rel		250.000		<input checked="" type="checkbox"/>
K_m_Ca_cyt	K_m_Ca_cyt		$5 \cdot 10^{-4}$		<input checked="" type="checkbox"/>
K_m_Ca_ds	K_m_Ca_ds		0.010		<input checked="" type="checkbox"/>
PrecFrac	PrecFrac		0.000		<input type="checkbox"/>
ActFrac	ActFrac		0.002		<input type="checkbox"/>
ProdFrac	ProdFrac		0.285		<input type="checkbox"/>
SpeedRel	SpeedRel		0.000		<input type="checkbox"/>
gama_SR_SL	gama_SR_SL		2.500		<input checked="" type="checkbox"/>
gama_SR_IT	gama_SR_IT		2.500		<input checked="" type="checkbox"/>
Na_o	Na_o		140.000		<input checked="" type="checkbox"/>
Na_i	Na_i		7.332		<input type="checkbox"/>
Ca_o	Ca_o		2.000		<input checked="" type="checkbox"/>
K_o	K_o		4.000		<input checked="" type="checkbox"/>
K_i	K_i		136.564		<input type="checkbox"/>
Ca_i-	Ca_i		$1.4 \cdot 10^{-5}$		<input type="checkbox"/>
_intracellular- _calcium- _concentration					
Ca_ds-	Ca_ds		$1.88 \cdot 10^{-5}$		<input type="checkbox"/>
_intracellular- _calcium- _concentration					
Ca_up	Ca_up		0.453		<input type="checkbox"/>
Ca_rel	Ca_rel		0.448		<input type="checkbox"/>
Ca_Calmod	Ca_Calmod		$5.555 \cdot 10^{-4}$		<input type="checkbox"/>
Ca_Trop	Ca_Trop		$2 \cdot 10^{-4}$		<input type="checkbox"/>
Calmod	Calmod		0.020		<input checked="" type="checkbox"/>
Trop	Trop		0.050		<input checked="" type="checkbox"/>
alpha_Calmod	alpha_Calmod		100000.000		<input checked="" type="checkbox"/>
beta_Calmod	beta_Calmod		50.000		<input checked="" type="checkbox"/>
alpha_Trop	alpha_Trop		0.000		<input type="checkbox"/>
beta_Trop	beta_Trop		200.000		<input checked="" type="checkbox"/>
gama_Trop_SL	gama_Trop_SL		1.500		<input checked="" type="checkbox"/>
KTrop	KTrop		5000.000		<input checked="" type="checkbox"/>
radius	radius		12.000		<input checked="" type="checkbox"/>
length	length		74.000		<input checked="" type="checkbox"/>
V_i	V_i		0.000		<input type="checkbox"/>

Id	Name	SBO	Value	Unit	Constant
V_Cell	V_Cell		0.000		<input type="checkbox"/>
V_i_ratio	V_i_ratio		0.000		<input type="checkbox"/>
V_ds_ratio	V_ds_ratio		0.100		<input checked="" type="checkbox"/>
V_rel_ratio	V_rel_ratio		0.100		<input checked="" type="checkbox"/>
V_e_ratio	V_e_ratio		0.400		<input checked="" type="checkbox"/>
V_up_ratio	V_up_ratio		0.010		<input checked="" type="checkbox"/>
Kdecay	Kdecay		10.000		<input checked="" type="checkbox"/>
light_chain	light_chain		$3.32 \cdot 10^{-5}$		<input type="checkbox"/>
cross_bridge	cross_bridge		$8.09 \cdot 10^{-5}$		<input type="checkbox"/>
KCont1	KCont1		12000.000		<input checked="" type="checkbox"/>
XCont2	XCont2		0.000		<input type="checkbox"/>
XCont1	XCont1		0.000		<input type="checkbox"/>
KCont2	KCont2		100.000		<input checked="" type="checkbox"/>
KCont3	KCont3		60.000		<input checked="" type="checkbox"/>
KCont4	KCont4		25.000		<input checked="" type="checkbox"/>
sarcomere- _length	sarcomere_length		2.000		<input checked="" type="checkbox"/>
cross- _bridge- _density	cross_bridge- _density		0.050		<input checked="" type="checkbox"/>
tension_rest	tension_rest		0.000		<input type="checkbox"/>
tension- _active	tension_active		0.000		<input type="checkbox"/>
overlap	overlap		0.000		<input type="checkbox"/>
cross- _bridge- _availability	cross_bridge- _availability		0.000		<input type="checkbox"/>
isometric- _tension	isometric_tension		0.000		<input type="checkbox"/>
i_stretch	i_stretch		0.000		<input type="checkbox"/>
i_Ca_stretch	i_Ca_stretch		0.000		<input type="checkbox"/>
i_K_stretch	i_K_stretch		0.000		<input type="checkbox"/>
i_Na_stretch	i_Na_stretch		0.000		<input type="checkbox"/>
i_An_stretch	i_An_stretch		0.000		<input type="checkbox"/>
gama_SAC_SL	gama_SAC_SL		2.500		<input checked="" type="checkbox"/>
gama_SAC_IT	gama_SAC_IT		2.500		<input checked="" type="checkbox"/>
SLHST	SLHST		2.000		<input checked="" type="checkbox"/>
ITHST	ITHST		1.000		<input checked="" type="checkbox"/>
g_Ca_stretch	g_Ca_stretch		0.010		<input checked="" type="checkbox"/>
g_K_stretch	g_K_stretch		0.010		<input checked="" type="checkbox"/>
g_Na_stretch	g_Na_stretch		0.010		<input checked="" type="checkbox"/>
g_An_stretch	g_An_stretch		0.010		<input checked="" type="checkbox"/>

Id	Name	SBO	Value	Unit	Constant
E_An_stretch	E_An_stretch		-20.000		<input checked="" type="checkbox"/>
f_stretch	f_stretch		0.000		<input type="checkbox"/>
g_fibro- _junct	g_fibro_junct		$2.9 \cdot 10^{-4}$		<input checked="" type="checkbox"/>
g_fibro	g_fibro		$2 \cdot 10^{-4}$		<input checked="" type="checkbox"/>
c_fibro	c_fibro		10^{-5}		<input checked="" type="checkbox"/>
g_fibro- _stretch	g_fibro_stretch		0.000		<input checked="" type="checkbox"/>
E_fibro- _stretch	E_fibro_stretch		0.000		<input checked="" type="checkbox"/>
V_fibro	V_fibro		-20.000		<input type="checkbox"/>
i_fibro	i_fibro		0.000		<input type="checkbox"/>
i_fibro- _junct	i_fibro_junct		0.000		<input type="checkbox"/>

5 Rules

This is an overview of 106 rules.

5.1 Rule `V_membrane`

Rule `V_membrane` is a rate rule for parameter `V_membrane`:

$$\frac{d}{dt}V_{\text{membrane}} = \frac{1}{C_m} \cdot (i_{\text{Stim}} + i_{\text{K1}} + i_{\text{to}} + i_{\text{Kr}} + i_{\text{Ks}} + i_{\text{K_ATP}} + i_{\text{K_ACh}} + i_{\text{NaK}} + i_{\text{Na}} + i_{\text{b_Na}} + i_{\text{p_Na}} + i_{\text{Ca_L_Na_cyt}} + i_{\text{Ca_L_Na_ds}} + i_{\text{NaCa_cyt}} + i_{\text{NaCa_ds}} + i_{\text{Ca_L_Ca_cyt}} + i_{\text{Ca_L_Ca_ds}} + i_{\text{Ca_L_K_cyt}} + i_{\text{Ca_L_K_ds}} + i_{\text{b_Ca}} + i_{\text{stretch}})$$

(1)

5.2 Rule `xr1`

Rule `xr1` is a rate rule for parameter `xr1`:

$$\frac{d}{dt}xr1 = \alpha_{xr1} \cdot (1 - xr1) - \beta_{xr1} \cdot xr1$$

(2)

5.3 Rule `xr2`

Rule `xr2` is a rate rule for parameter `xr2`:

$$\frac{d}{dt}xr2 = \alpha_{xr2} \cdot (1 - xr2) - \beta_{xr2} \cdot xr2$$

(3)

5.4 Rule xs

Rule xs is a rate rule for parameter xs :

$$\frac{d}{dt}xs = \text{alpha_xs} \cdot (1 - xs) - \text{beta_xs} \cdot xs \quad (4)$$

5.5 Rule m

Rule m is a rate rule for parameter m :

$$\frac{d}{dt}m = \text{alpha_m} \cdot (1 - m) - \text{beta_m} \cdot m \quad (5)$$

5.6 Rule h

Rule h is a rate rule for parameter h :

$$\frac{d}{dt}h = \text{alpha_h} \cdot (1 - h) - \text{beta_h} \cdot h \quad (6)$$

5.7 Rule d

Rule d is a rate rule for parameter d :

$$\frac{d}{dt}d = \text{speed_d} \cdot (\text{alpha_d} \cdot (1 - d) - \text{beta_d} \cdot d) \quad (7)$$

5.8 Rule f

Rule f is a rate rule for parameter f :

$$\frac{d}{dt}f = \text{speed_f} \cdot (\text{alpha_f} \cdot (1 - f) - \text{beta_f} \cdot f) \quad (8)$$

5.9 Rule $f2$

Rule $f2$ is a rate rule for parameter $f2$:

$$\frac{d}{dt}f2 = 1 - 1 \cdot \left(\frac{\text{Ca_i_L_type_Ca_channel_f2_gate}}{\text{Km_f2} + \text{Ca_i_L_type_Ca_channel_f2_gate}} + f2 \right) \quad (9)$$

5.10 Rule $f2ds$

Rule $f2ds$ is a rate rule for parameter $f2ds$:

$$\frac{d}{dt}f2ds = \text{R_decay} \cdot \left(1 - \left(\frac{\text{Ca_ds_L_type_Ca_channel_f2ds_gate}}{\text{Km_f2ds} + \text{Ca_ds_L_type_Ca_channel_f2ds_gate}} + f2ds \right) \right) \quad (10)$$

5.11 Rule s

Rule s is a rate rule for parameter s :

$$\frac{d}{dt}s = \text{alpha}_s \cdot (1 - s) - \text{beta}_s \cdot s \quad (11)$$

5.12 Rule r

Rule r is a rate rule for parameter r :

$$\frac{d}{dt}r = 333 \cdot \left(\frac{1}{1 + \exp\left(\frac{(\text{V_transient_outward_current_r_gate} + 4)}{5}\right)} - r \right) \quad (12)$$

5.13 Rule x_ACh

Rule x_ACh is a rate rule for parameter x_ACh :

$$\frac{d}{dt}x_ACh = \text{alpha_ACh} \cdot (1 - x_ACh) - \text{beta_ACh} \cdot x_ACh \quad (13)$$

5.14 Rule $ActFrac$

Rule $ActFrac$ is a rate rule for parameter $ActFrac$:

$$\frac{d}{dt}ActFrac = \text{PrecFrac} \cdot \text{SpeedRel} \cdot \text{ActRate} - ActFrac \cdot \text{SpeedRel} \cdot \text{InactRate} \quad (14)$$

5.15 Rule $ProdFrac$

Rule $ProdFrac$ is a rate rule for parameter $ProdFrac$:

$$\frac{d}{dt}ProdFrac = ActFrac \cdot \text{SpeedRel} \cdot \text{InactRate} - \text{SpeedRel} \cdot 1 \cdot ProdFrac \quad (15)$$

5.16 Rule Na_i

Rule Na_i is a rate rule for parameter Na_i :

$$\frac{d}{dt}Na_i = \frac{1}{1 \cdot V_i \cdot F} \cdot (i_Na + i_p_Na + i_b_Na + 3 \cdot i_NaK + 3 \cdot i_NaCa_cyt + i_Ca_L_Na_cyt + i_Ca_L_Na_ds) \quad (16)$$

5.17 Rule K_i

Rule K_i is a rate rule for parameter K_i :

$$\frac{d}{dt}K_i = \frac{1}{1 \cdot V_i \cdot F} \cdot (i_K1 + i_Kr + i_Ks + i_Ca_L_K_cyt + i_Ca_L_K_ds + i_to - 2 \cdot i_NaK) \quad (17)$$

5.18 Rule Ca_i_intracellular_calcium_concentration

Rule Ca_i_intracellular_calcium_concentration is a rate rule for parameter Ca_i_intracellular_calcium_concentration:

$$\begin{aligned} \frac{d}{dt} \text{Ca}_i \text{intracellular_calcium_concentration} = & \frac{1}{2 \cdot 1 \cdot V_i \cdot F} \cdot (i_{Ca_L_Ca_cyt} + i_{b_Ca} - 2 \\ & \cdot i_{NaCa_cyt} - 2 \cdot i_{NaCa_ds}) \\ & + \text{Ca}_{ds} \text{intracellular_calcium_concentration} \\ & \cdot V_{ds_ratio} \cdot K_{decay} + \frac{i_{rel} \cdot V_{rel_ratio}}{V_i \text{ratio}} \\ & - \text{Ca_Calmod} - \text{Ca_Trop} - i_{up} \end{aligned} \quad (18)$$

5.19 Rule Ca_ds_intracellular_calcium_concentration

Rule Ca_ds_intracellular_calcium_concentration is a rate rule for parameter Ca_ds_intracellular_calcium_concentration:

$$\begin{aligned} \frac{d}{dt} \text{Ca}_{ds} \text{intracellular_calcium_concentration} = & \frac{1 \cdot i_{Ca_L_Ca_ds}}{2 \cdot 1 \cdot V_{ds_ratio} \cdot V_i \cdot F} \\ & - \text{Ca}_{ds} \text{intracellular_calcium_concentration} \\ & \cdot K_{decay} \end{aligned} \quad (19)$$

5.20 Rule Ca_up

Rule Ca_up is a rate rule for parameter Ca_up:

$$\frac{d}{dt} \text{Ca}_{up} = \frac{V_i \text{ratio}}{V_{up_ratio}} \cdot i_{up} - i_{trans} \quad (20)$$

5.21 Rule Ca_rel

Rule Ca_rel is a rate rule for parameter Ca_rel:

$$\frac{d}{dt} \text{Ca}_{rel} = \frac{V_{up_ratio}}{V_{rel_ratio}} \cdot i_{trans} - i_{rel} \quad (21)$$

5.22 Rule Ca_Calmod

Rule Ca_Calmod is a rate rule for parameter Ca_Calmod:

$$\begin{aligned} \frac{d}{dt} \text{Ca_Calmod} = & \alpha_{Calmod} \cdot \text{Ca}_i \text{intracellular_calcium_concentration} \\ & \cdot (\text{Calmod} - \text{Ca_Calmod}) - \beta_{Calmod} \cdot \text{Ca_Calmod} \end{aligned} \quad (22)$$

5.23 Rule Ca_Trop

Rule Ca_Trop is a rate rule for parameter Ca_Trop:

$$\frac{d}{dt} \text{Ca_Trop} = \text{alpha_Trop} \cdot \text{Ca_i_intracellular_calcium_concentration} \cdot (\text{Trop} - \text{Ca_Trop}) - \text{beta_Trop} \cdot \text{Ca_Trop} \quad (23)$$

5.24 Rule light_chain

Rule light_chain is a rate rule for parameter light_chain:

$$\frac{d}{dt} \text{light_chain} = \text{KCont1} \cdot \text{XCont1}^2 \cdot \text{XCont2} \cdot (1 - \text{light_chain}) - \text{KCont2} \cdot \text{light_chain} \quad (24)$$

5.25 Rule cross_bridge

Rule cross_bridge is a rate rule for parameter cross_bridge:

$$\frac{d}{dt} \text{cross_bridge} = \text{KCont3} \cdot \text{light_chain} \cdot (1 - \text{cross_bridge}) - \text{KCont4} \cdot \text{cross_bridge} \quad (25)$$

5.26 Rule V_fibro

Rule V_fibro is a rate rule for parameter V_fibro:

$$\frac{d}{dt} \text{V_fibro} = \frac{(\text{i_fibro} + \text{i_fibro_junct})}{\text{c_fibro}} \quad (26)$$

5.27 Rule i_Stim

Rule i_Stim is an assignment rule for parameter i_Stim:

$$\text{i_Stim} = \begin{cases} \text{stim_amplitude} & \text{if } (\text{time} \geq \text{stim_start}) \wedge \left(\text{time} - \text{stim_start} - \left\lfloor \frac{\text{time} - \text{stim_start}}{\text{stim_period}} \right\rfloor \cdot \text{stim_period} \leq \text{stim_duration} \right) \\ 0 & \text{otherwise} \end{cases} \quad (27)$$

5.28 Rule E_Na

Rule E_Na is an assignment rule for parameter E_Na:

$$\text{E_Na} = \frac{R \cdot T}{F} \cdot \left(\frac{\text{Na_o}}{\text{Na_i}} \right) \quad (28)$$

5.29 Rule E_K

Rule E_K is an assignment rule for parameter E_K:

$$\text{E_K} = \frac{R \cdot T}{F} \cdot \left(\frac{\text{K_o}}{\text{K_i}} \right) \quad (29)$$

5.30 Rule E_Ks

Rule E_Ks is an assignment rule for parameter E_Ks:

$$E_Ks = \frac{R \cdot T}{F} \cdot \left(\frac{K_o + P_kna \cdot Na_o}{K_i + P_kna \cdot Na_i} \right) \quad (30)$$

5.31 Rule E_Ca

Rule E_Ca is an assignment rule for parameter E_Ca:

$$E_Ca = \frac{0.5 \cdot R \cdot T}{F} \cdot \left(\frac{Ca_o}{Ca_i \text{ intracellular_calcium_concentration}} \right) \quad (31)$$

5.32 Rule E_mh

Rule E_mh is an assignment rule for parameter E_mh:

$$E_mh = \frac{R \cdot T}{F} \cdot \left(\frac{Na_o + 0.12 \cdot K_o}{Na_i + 0.12 \cdot K_i} \right) \quad (32)$$

5.33 Rule i_K1

Rule i_K1 is an assignment rule for parameter i_K1:

$$i_K1 = \frac{\frac{g_K1 \cdot K_o}{K_o + K_mk1} \cdot (V_membrane - E_K)}{1 + \exp\left(\frac{(V_membrane - E_K - 10) \cdot F \cdot 1.25}{R \cdot T}\right)} \quad (33)$$

5.34 Rule i_Kr

Rule i_Kr is an assignment rule for parameter i_Kr:

$$i_Kr = \frac{(g_Kr1 \cdot xr1 + g_Kr2 \cdot xr2) \cdot 1}{1 + \exp\left(\frac{V_membrane + 9}{22.4}\right)} \cdot (V_membrane - E_K) \quad (34)$$

5.35 Rule alpha_xr1

Rule alpha_xr1 is an assignment rule for parameter alpha_xr1:

$$\alpha_xr1 = \frac{50}{1 + \exp\left(\frac{(V_rapid_delayed_rectifier_potassium_current_xr1_gate - 5)}{9}\right)} \quad (35)$$

5.36 Rule beta_xr1

Rule beta_xr1 is an assignment rule for parameter beta_xr1:

$$\beta_xr1 = 0.05 \cdot \exp\left(\frac{(V_rapid_delayed_rectifier_potassium_current_xr1_gate - 20)}{15}\right) \quad (36)$$

5.37 Rule `alpha_xr2`

Rule `alpha_xr2` is an assignment rule for parameter `alpha_xr2`:

$$\text{alpha_xr2} = \frac{50}{1 + \exp\left(\frac{(\text{V_rapid_delayed_rectifier_potassium_current_xr2_gate} - 5)}{9}\right)} \quad (37)$$

5.38 Rule `beta_xr2`

Rule `beta_xr2` is an assignment rule for parameter `beta_xr2`:

$$\text{beta_xr2} = 0.4 \cdot \exp\left(\left(\left(\frac{\text{V_rapid_delayed_rectifier_potassium_current_xr2_gate} + 30}{30}\right)^3\right)\right) \quad (38)$$

5.39 Rule `i_Ks`

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

$$i_Ks = g_Ks \cdot xs^2 \cdot (\text{V_membrane} - E_Ks) \quad (39)$$

5.40 Rule `alpha_xs`

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

$$\text{alpha_xs} = \frac{14}{1 + \exp\left(\frac{(\text{V_slow_delayed_rectifier_potassium_current_xs_gate} - 40)}{9}\right)} \quad (40)$$

5.41 Rule `beta_xs`

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

$$\text{beta_xs} = 1 \cdot \exp\left(\frac{\text{V_slow_delayed_rectifier_potassium_current_xs_gate}}{45}\right) \quad (41)$$

5.42 Rule `i_K_ATP`

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

$$i_K_ATP = \frac{g_K_ATP \cdot (\text{V_membrane} + 80)}{1 + \left(\frac{ATP}{K_ATP}\right)^2} \quad (42)$$

5.43 Rule `i_KNa`

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

$$i_KNa = \frac{g_K_Na \cdot Na_i}{Na_i + K_kna} \cdot (\text{V_membrane} - E_K) \quad (43)$$

5.44 Rule i_{Na}

Rule i_{Na} is an assignment rule for parameter i_{Na} :

$$i_{Na} = g_{Na} \cdot m^3 \cdot h \cdot (V_{membrane} - E_{mh}) \quad (44)$$

5.45 Rule $E0_m$

Rule $E0_m$ is an assignment rule for parameter $E0_m$:

$$E0_m = V_{fast_sodium_current_m_gate} + 41 \quad (45)$$

5.46 Rule α_m

Rule α_m is an assignment rule for parameter α_m :

$$\alpha_m = \begin{cases} 2000 & \text{if } |E0_m| < \delta_{\alpha_m} \\ \frac{200 \cdot E0_m}{1 - \exp(0.1 \cdot E0_m)} & \text{otherwise} \end{cases} \quad (46)$$

5.47 Rule β_m

Rule β_m is an assignment rule for parameter β_m :

$$\beta_m = 8000 \cdot \exp(0.056 \cdot (V_{fast_sodium_current_m_gate} + 66)) \quad (47)$$

5.48 Rule α_h

Rule α_h is an assignment rule for parameter α_h :

$$\alpha_h = 20 \cdot \exp(0.125 \cdot (V_{fast_sodium_current_h_gate} + 75)) \quad (48)$$

5.49 Rule β_h

Rule β_h is an assignment rule for parameter β_h :

$$\beta_h = \frac{2000}{1 + 320 \cdot \exp(0.1 \cdot (V_{fast_sodium_current_h_gate} + 75))} \quad (49)$$

5.50 Rule i_{p_Na}

Rule i_{p_Na} is an assignment rule for parameter i_{p_Na} :

$$i_{p_Na} = \frac{g_{pna} \cdot 1}{1 + \exp\left(\frac{(V_{membrane} + 52)}{8}\right)} \cdot (V_{membrane} - E_{Na}) \quad (50)$$

5.51 Rule i_b_Na

Rule i_b_Na is an assignment rule for parameter i_b_Na :

$$i_b_Na = g_bna \cdot (V_membrane - E_Na) \quad (51)$$

5.52 Rule $i_Ca_L_Ca_cyt$

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

$$i_Ca_L_Ca_cyt = \frac{(1-FrCa) \cdot 4 \cdot P_Ca_L \cdot d \cdot f \cdot f2 \cdot (V_membrane - 50) \cdot F}{R \cdot T} \cdot \left(Ca_i_intracellular_calcium_concentration \cdot \exp\left(\frac{100 \cdot F}{R \cdot T}\right) - Ca_o \right) \cdot \exp\left(\frac{(V_membrane - 50) \cdot F \cdot 2}{R \cdot T}\right) \quad (52)$$

5.53 Rule $i_Ca_L_K_cyt$

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

$$i_Ca_L_K_cyt = \frac{(1-FrCa) \cdot P_CaK \cdot P_Ca_L \cdot d \cdot f \cdot f2 \cdot (V_membrane - 50) \cdot F}{R \cdot T} \cdot \left(K_i \cdot \exp\left(\frac{50 \cdot F}{R \cdot T}\right) - K_o \cdot \exp\left(\frac{(V_membrane - 50) \cdot F}{R \cdot T}\right) \right) \quad (53)$$

5.54 Rule $i_Ca_L_Na_cyt$

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

$$i_Ca_L_Na_cyt = \frac{(1-FrCa) \cdot P_CaNa \cdot P_Ca_L \cdot d \cdot f \cdot f2 \cdot (V_membrane - 50) \cdot F}{R \cdot T} \cdot \left(Na_i \cdot \exp\left(\frac{50 \cdot F}{R \cdot T}\right) - Na_o \cdot \exp\left(\frac{(V_membrane - 50) \cdot F}{R \cdot T}\right) \right) \quad (54)$$

5.55 Rule $i_Ca_L_Ca_ds$

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

$$i_Ca_L_Ca_ds = \frac{FrCa \cdot 4 \cdot P_Ca_L \cdot d \cdot f \cdot f2 \cdot (V_membrane - 50) \cdot F}{R \cdot T} \cdot \left(Ca_i_intracellular_calcium_concentration \cdot \exp\left(\frac{100 \cdot F}{R \cdot T}\right) - Ca_o \cdot \exp\left(\frac{(V_membrane - 50) \cdot F \cdot 2}{R \cdot T}\right) \right) \quad (55)$$

5.56 Rule $i_Ca_L_K_ds$

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

$$i_Ca_L_K_ds = \frac{\text{FriCa}_P\text{-CaK}_P\text{-Ca}_L\text{-d}_f\text{f2ds}\cdot(V_membrane-50)\cdot F}{R\cdot T} \cdot \left(K_i \cdot \exp\left(\frac{50\cdot F}{R\cdot T}\right) - K_o \cdot \exp\left(\frac{(V_membrane-50)\cdot F}{R\cdot T}\right) \right) \quad (56)$$

5.57 Rule $i_Ca_L_Na_ds$

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

$$i_Ca_L_Na_ds = \frac{\text{FriCa}_P\text{-CaNa}_P\text{-Ca}_L\text{-d}_f\text{f2ds}\cdot(V_membrane-50)\cdot F}{R\cdot T} \cdot \left(Na_i \cdot \exp\left(\frac{50\cdot F}{R\cdot T}\right) - Na_o \cdot \exp\left(\frac{(V_membrane-50)\cdot F}{R\cdot T}\right) \right) \quad (57)$$

5.58 Rule i_Ca_L

Rule i_Ca_L is an assignment rule for parameter i_Ca_L :

$$i_Ca_L = i_Ca_L_Ca_cyt + i_Ca_L_K_cyt + i_Ca_L_Na_cyt + i_Ca_L_Ca_ds + i_Ca_L_K_ds + i_Ca_L_Na_ds \quad (58)$$

5.59 Rule $E0_d$

Rule $E0_d$ is an assignment rule for parameter $E0_d$:

$$E0_d = V_L.type_Ca_channel.d_gate + 24 - 5 \quad (59)$$

5.60 Rule α_d

Rule α_d is an assignment rule for parameter α_d :

$$\alpha_d = \begin{cases} 120 & \text{if } |E0_d| < 10^{-4} \\ \frac{30 \cdot E0_d}{1 - \exp\left(\frac{E0_d}{4}\right)} & \text{otherwise} \end{cases} \quad (60)$$

5.61 Rule β_d

Rule β_d is an assignment rule for parameter β_d :

$$\beta_d = \begin{cases} 120 & \text{if } |E0_d| < 10^{-4} \\ \frac{12 \cdot E0_d}{\exp\left(\frac{E0_d}{10}\right) - 1} & \text{otherwise} \end{cases} \quad (61)$$

5.62 Rule $E0_f$

Rule $E0_f$ is an assignment rule for parameter $E0_f$:

$$E0_f = V_L_type_Ca_channel_f_gate + 34 \quad (62)$$

5.63 Rule $alpha_f$

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

$$alpha_f = \begin{cases} 25 & \text{if } |E0_f| < delta_f \\ \frac{6.25 \cdot E0_f}{\exp\left(\frac{E0_f}{4}\right) - 1} & \text{otherwise} \end{cases} \quad (63)$$

5.64 Rule $beta_f$

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

$$beta_f = \frac{12}{1 + \exp\left(\frac{1 \cdot (V_L_type_Ca_channel_f_gate + 34)}{4}\right)} \quad (64)$$

5.65 Rule i_b_Ca

Rule i_b_Ca is an assignment rule for parameter i_b_Ca :

$$i_b_Ca = g_bca \cdot (V_membrane - E_Ca) \quad (65)$$

5.66 Rule i_to

Rule i_to is an assignment rule for parameter i_to :

$$i_to = g_to \cdot (g_tos + s \cdot (1 - g_tos)) \cdot r \cdot (V_membrane - E_K) \quad (66)$$

5.67 Rule $alpha_s$

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

$$alpha_s = 0.033 \cdot \exp\left(\frac{V_transient_outward_current_s_gate}{17}\right) \quad (67)$$

5.68 Rule $beta_s$

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

$$beta_s = \frac{33}{1 + \exp(0.125 \cdot (V_transient_outward_current_s_gate + 10))} \quad (68)$$

5.69 Rule i_K_ACh

Rule i_K_ACh is an assignment rule for parameter i_K_ACh :

$$i_K_ACh = \frac{g_KACh \cdot \frac{K_o}{K_o + K_{mkI}} \cdot x_ACh \cdot \frac{1}{1 + \left(\frac{K_D}{ACh}\right)^2} \cdot (V_membrane - E_K)}{1 + \exp\left(\frac{2 \cdot F \cdot (V_membrane - (E_K + 10))}{R \cdot T}\right)} \quad (69)$$

5.70 Rule i_NaK

Rule i_NaK is an assignment rule for parameter i_NaK :

$$i_NaK = \frac{i_NaK_max \cdot K_o}{K_{mK} + K_o} \cdot Na_i \quad (70)$$

5.71 Rule i_NaCa_cyt

Rule i_NaCa_cyt is an assignment rule for parameter i_NaCa_cyt :

$$i_NaCa_cyt = \frac{(1 - FRiNaCa) \cdot k_NaCa \cdot \left(\exp\left(\frac{\gamma \cdot (n_NaCa - 2) \cdot V_membrane \cdot F}{R \cdot T}\right) \cdot Na_i^{n_NaCa} \cdot Ca_o - \exp\left(\frac{(\gamma - 1) \cdot (n_NaCa - 2) \cdot V_membrane \cdot F}{R \cdot T}\right) \cdot Na_o^{n_NaCa} \cdot Ca_o\right)}{(1 + d_NaCa \cdot (Ca_i_intracellular_calcium_concentration \cdot Na_o^{n_NaCa} + Ca_o \cdot Na_i^{n_NaCa}))} \quad (71)$$

5.72 Rule i_NaCa_ds

Rule i_NaCa_ds is an assignment rule for parameter i_NaCa_ds :

$$i_NaCa_ds = \frac{FRiNaCa \cdot k_NaCa \cdot \left(\exp\left(\frac{\gamma \cdot (n_NaCa - 2) \cdot V_membrane \cdot F}{R \cdot T}\right) \cdot Na_i^{n_NaCa} \cdot Ca_o - \exp\left(\frac{(\gamma - 1) \cdot (n_NaCa - 2) \cdot V_membrane \cdot F}{R \cdot T}\right) \cdot Na_o^{n_NaCa} \cdot Ca_o\right)}{(1 + d_NaCa \cdot (Ca_ds_intracellular_calcium_concentration \cdot Na_o^{n_NaCa} + Ca_o \cdot Na_i^{n_NaCa}))} \quad (72)$$

5.73 Rule i_NaCa

Rule i_NaCa is an assignment rule for parameter i_NaCa :

$$i_NaCa = i_NaCa_cyt + i_NaCa_ds \quad (73)$$

5.74 Rule K_1

Rule K_1 is an assignment rule for parameter K_1 :

$$K_1 = \frac{K_cyca \cdot K_xcs}{K_srca} \quad (74)$$

5.75 Rule K_2

Rule K_2 is an assignment rule for parameter K_2 :

$$K_2 = Ca_i_intracellular_calcium_concentration + Ca_up \cdot K_1 + K_cyca \cdot K_xcs + K_cyca \quad (75)$$

5.76 Rule `i_up`

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

$$i_up = \frac{Ca_i_intracellular_calcium_concentration}{K_2} \cdot alpha_up - \frac{Ca_up \cdot K_1}{K_2} \cdot beta_up \quad (76)$$

5.77 Rule `i_trans`

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

$$i_trans = 50 \cdot (Ca_up - Ca_rel) \quad (77)$$

5.78 Rule `VoltDep`

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

$$VoltDep = \exp(0.08 \cdot (V_membrane - 40)) \quad (78)$$

Derived unit dimensionless

5.79 Rule `CaiReg`

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

$$CaiReg = \frac{Ca_i_intracellular_calcium_concentration}{Ca_i_intracellular_calcium_concentration + K_m_Ca_cyt} \quad (79)$$

5.80 Rule `CadsReg`

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

$$CadsReg = \frac{Ca_ds_intracellular_calcium_concentration}{Ca_ds_intracellular_calcium_concentration + K_m_Ca_ds} \quad (80)$$

5.81 Rule `RegBindSite`

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

$$RegBindSite = CaiReg + (1 - CaiReg) \cdot CadsReg \quad (81)$$

5.82 Rule `ActRate`

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

$$ActRate = 0 \cdot VoltDep + 500 \cdot RegBindSite^2 \quad (82)$$

5.83 Rule `InactRate`

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

$$\text{InactRate} = 60 + 500 \cdot \text{RegBindSite}^2 \quad (83)$$

5.84 Rule `SpeedRel`

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

$$\text{SpeedRel} = \begin{cases} 5 & \text{if } V_{\text{membrane}} < 50 \\ 1 & \text{otherwise} \end{cases} \quad (84)$$

5.85 Rule `SRLeak`

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

$$\text{SRLeak} = \begin{cases} K_{\text{leak_rate}} \cdot \exp(\text{gama_SR_IT} \cdot \text{isometric_tension}) & \text{if } \text{isometric_tension} > 0 \\ K_{\text{leak_rate}} \cdot \exp(\text{gama_SR_SL} \cdot \text{sarcomere_length}) & \text{otherwise} \end{cases} \quad (85)$$

5.86 Rule `PrecFrac`

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

$$\text{PrecFrac} = 1 - \text{ActFrac} - \text{ProdFrac} \quad (86)$$

5.87 Rule `i_rel`

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

$$i_{\text{rel}} = \left(\left(\frac{\text{ActFrac}}{\text{ActFrac} + 0.25} \right)^2 \cdot K_{\text{m_rel}} + \text{SRLeak} \right) \cdot \text{Ca}_{\text{rel}} \quad (87)$$

5.88 Rule `V_Cell`

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

$$V_{\text{Cell}} = \frac{3.141592654 \cdot \left(\frac{\text{radius}}{1000} \right)^2 \cdot \text{length}}{1000} \quad (88)$$

5.89 Rule `V_i_ratio`

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

$$V_{\text{i_ratio}} = 1 - V_{\text{e_ratio}} - V_{\text{up_ratio}} - V_{\text{rel_ratio}} \quad (89)$$

5.90 Rule `V_i`

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

$$V_i = V_{Cell} \cdot V_{i_ratio} \quad (90)$$

5.91 Rule `alpha_Trop`

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

$$\alpha_{Trop} = K_{Trop} \cdot \exp(\gamma_{Trop_SL} \cdot sarcomere_length) \quad (91)$$

5.92 Rule `XCont2`

Rule `XCont2` is an assignment rule for parameter `XCont2`:

$$X_{Cont2} = \frac{Ca_Calmod}{Calmod} \quad (92)$$

5.93 Rule `XCont1`

Rule `XCont1` is an assignment rule for parameter `XCont1`:

$$X_{Cont1} = \frac{Ca_Trop}{Trop} \quad (93)$$

5.94 Rule `tension_rest`

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

$$tension_rest = 2 \cdot 10^{-4} \cdot \exp(2 \cdot sarcomere_length) \quad (94)$$

5.95 Rule `tension_active`

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

$$tension_active = \begin{cases} 1 - \exp(3 \cdot (sarcomere_length - 1)) & \text{if } sarcomere_length > 1 \\ 0 & \text{otherwise} \end{cases} \quad (95)$$

5.96 Rule `overlap`

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

$$overlap = \begin{cases} 1 - 0.625 \cdot (sarcomere_length - 2) & \text{if } sarcomere_length > 2 \\ 1 & \text{otherwise} \end{cases} \quad (96)$$

5.97 Rule `cross_bridge_availability`

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

$$\text{cross_bridge_availability} = \text{tension_active} \cdot \text{overlap} \cdot \text{cross_bridge_density} \quad (97)$$

5.98 Rule `isometric_tension`

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

$$\text{isometric_tension} = \text{cross_bridge} \cdot \text{cross_bridge_availability} + \text{tension_rest} \quad (98)$$

5.99 Rule `f_stretch`

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

$$f_stretch = \begin{cases} \frac{1}{1 + \exp(2 \cdot \text{gama_SAC_IT} \cdot (\text{isometric_tension} - \text{ITHST}))} & \text{if isometric_tension} > 0 \\ \frac{1}{1 + \exp(2 \cdot \text{gama_SAC_SL} \cdot (\text{sarcomere_length} - \text{SLHST}))} & \text{otherwise} \end{cases} \quad (99)$$

5.100 Rule `i_Ca_stretch`

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

$$i_Ca_stretch = g_Ca_stretch \cdot f_stretch \cdot (V_membrane - E_Ca) \quad (100)$$

5.101 Rule `i_Na_stretch`

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

$$i_Na_stretch = g_Na_stretch \cdot f_stretch \cdot (V_membrane - E_Na) \quad (101)$$

5.102 Rule `i_K_stretch`

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

$$i_K_stretch = g_K_stretch \cdot f_stretch \cdot (V_membrane - E_K) \quad (102)$$

5.103 Rule `i_An_stretch`

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

$$i_An_stretch = g_An_stretch \cdot f_stretch \cdot (V_membrane - E_An_stretch) \quad (103)$$

5.104 Rule `i_stretch`

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

$$i_stretch = i_Ca_stretch + i_Na_stretch + i_K_stretch + i_An_stretch \quad (104)$$

5.105 Rule `i_fibro`

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

$$i_fibro = g_fibro \cdot (V_fibro + 20) + g_fibro_stretch \cdot (V_fibro - E_fibro_stretch) \quad (105)$$

5.106 Rule `i_fibro_junct`

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

$$i_fibro_junct = g_fibro_junct \cdot (V_membrane - V_fibro) \quad (106)$$

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