# **SBML Model Report**

# Model name: "Izhikevich2004-\_SpikingNeurons\_thresholdVariability"



May 5, 2016

#### 1 General Overview

This is a document in SBML Level 2 Version 4 format. This model was created by Enuo He<sup>1</sup> at July 16<sup>th</sup> 2007 at 9:41 a. m. and last time modified at February 25<sup>th</sup> 2015 at 11:18 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	Quantity   Element	
compartment types	0	compartments	1
species types	0	species	0
events	4	constraints	0
reactions	0	function definitions	0
global parameters	8	unit definitions	0
rules	2	initial assignments	0

#### **Model Notes**

This a model from the article:

Which model to use for cortical spiking neurons?

Izhikevich EM. IEEE Trans Neural Netw.2004 Sep;15(5):1063-70. 15484883,

#### **Abstract:**

We discuss the biological plausibility and computational efficiency of some of the most useful

<sup>&</sup>lt;sup>1</sup>BNMC, enuo@caltech.edu

models of spiking and bursting neurons. We compare their applicability to large-scale simulations of cortical neural networks.

The model is according to the paper Which Model to Use for Cortical Spiking Neurons? Figure 1(O) threshold variability has been reproduced by MathSBML. The ODE and the parameters values are taken from the a paper Simple Model of Spiking Neurons The original format of the models are encoded in the MATLAB format existed in the ModelDB with Accession number 39948

Figure 1 are the simulation results of the same model with different choices of parameters and different stimulus function or events.a=0.03; b=0.25; c=-60; d=4; V=-64; u=b\*V;

This model originates from BioModels Database: A Database of Annotated Published Models. It is copyright (c) 2005-2010 The BioModels Team.

For more information see the terms of use.

To cite BioModels Database, please use Le Novre N., Bornstein B., Broicher A., Courtot M., Donizelli M., Dharuri H., Li L., Sauro H., Schilstra M., Shapiro B., Snoep J.L., Hucka M. (2006) BioModels Database: A Free, Centralized Database of Curated, Published, Quantitative Kinetic Models of Biochemical and Cellular Systems Nucleic Acids Res., 34: D689-D691.

#### 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<sup>2</sup>

#### 2.4 Unit length

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

**Definition** m

#### 2.5 Unit time

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

**Definition** s

# 3 Compartment

This model contains one compartment.

Table 2: Properties of all compartments.

Id	Name	SBO	Spatial Dimensions	Size	Unit	Constant	Outside
cell			3	1	litre	Ø	

# **3.1 Compartment** cell

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

# **4 Parameters**

This model contains eight global parameters.

Table 3: Properties of each parameter.

F F							
Id	Name	SBO Value	Unit	Constant			
a		0.03		$ \overline{\mathbf{Z}} $			
b		0.25					
С		-60.00					
d		4.00					
Vthresh		30.00					
i		0.00					
v		-64.00					
u		-16.00					

# 5 Rules

This is an overview of two rules.

#### 5.1 Rule **▽**

Rule v is a rate rule for parameter v:

$$\frac{d}{dt}v = 0.04 \cdot v^2 + 5 \cdot v + 140 - u + i \tag{1}$$

#### **5.2 Rule** u

Rule u is a rate rule for parameter u:

$$\frac{\mathrm{d}}{\mathrm{d}t}\mathbf{u} = \mathbf{a} \cdot (\mathbf{b} \cdot \mathbf{v} - \mathbf{u}) \tag{2}$$

# 6 Events

This is an overview of four events. Each event is initiated whenever its trigger condition switches from false to true. A delay function postpones the effects of an event to a later time point. At the time of execution, an event can assign values to species, parameters or compartments if these are not set to constant.

#### **6.1 Event** event\_0000001

# **Trigger condition**

$$v > V$$
thresh (3)

# **Assignments**

$$v = c \tag{4}$$

$$u = u + d \tag{5}$$

#### **6.2 Event** event\_0000002

#### **Trigger condition**

$$((time > 10) \land (time < 15)) \lor ((time > 80) \land (time < 85))$$

#### **Assignment**

$$i = 1 \tag{7}$$

# **6.3 Event** event\_0000003

# **Trigger condition**

$$(time > 70) \land (time < 75) \tag{8}$$

#### **Assignment**

$$i = -6 \tag{9}$$

# **6.4 Event** event\_0000004

#### **Trigger condition**

$$((time \ge 15) \land (time \le 70)) \lor (time \ge 85) \tag{10}$$

# **Assignment**

$$i = 0 \tag{11}$$

 $\mathfrak{BML}^{\underline{\mathsf{ATEX}}}$  was developed by Andreas Dräger<sup>a</sup>, Hannes Planatscher<sup>a</sup>, Dieudonné M Wouamba<sup>a</sup>, Adrian Schröder<sup>a</sup>, Michael Hucka<sup>b</sup>, Lukas Endler<sup>c</sup>, Martin Golebiewski<sup>d</sup> and Andreas Zell<sup>a</sup>. Please see http://www.ra.cs.uni-tuebingen.de/software/SBML2LaTeX for more information.

<sup>&</sup>lt;sup>a</sup>Center for Bioinformatics Tübingen (ZBIT), Germany

<sup>&</sup>lt;sup>b</sup>California Institute of Technology, Beckman Institute BNMC, Pasadena, United States

<sup>&</sup>lt;sup>c</sup>European Bioinformatics Institute, Wellcome Trust Genome Campus, Hinxton, United Kingdom

<sup>&</sup>lt;sup>d</sup>EML Research gGmbH, Heidelberg, Germany