



Cybow Modeller

modeller

- Yet Another Cell Modelling Environment

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Introduction

- Cybow Modeller is a software suite for cell modelling
 - Cybow [saibou] 細胞: “cell” in Japanese
- Features
 - Graphical & user-friendly interface with assists
 - Semantic annotations of variables with CPO
 - C++ code generator (under development)
 - OS independent: implemented in Java & XSLT
 - Open source: available at SourceForge.net

Modelling Workflow

Define components

IKATP.cmc - Cybow Component Editor

Component Name: KATP_current

Description: ATP-sensitive potassium channel current

Variable

R	Name	Symbol	Unit	Default	Concept
<input checked="" type="checkbox"/>	t	t	ms		time
<input type="checkbox"/>	IKATP	I _{KATP}	mA		ATP-sensitive_potassium_current
<input type="checkbox"/>	N	N		2.333E3	
<input type="checkbox"/>	gamma	γ			conductance of ATP-sensitive
<input checked="" type="checkbox"/>	Vm	V _m	mV		cell_membrane_potential
<input checked="" type="checkbox"/>	EK	E _K	mV		potassium_equilibrium_potential
<input type="checkbox"/>	p_open	p(open)			open probability of ATP-sensitive
<input checked="" type="checkbox"/>	K_o	[K ⁺] _o	mM		external_potassium_concentration
<input checked="" type="checkbox"/>	ATP_i	[ATP] _i	mM		intracellular_ATP_concentration

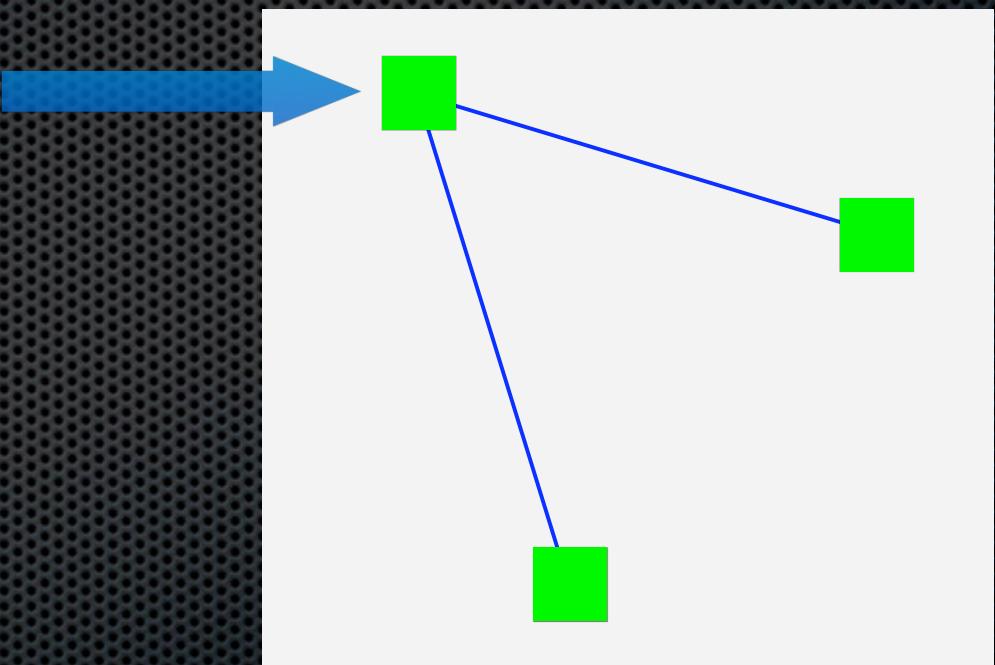
Equations

$$I_{KATP} = N \cdot \gamma \cdot (V_m - E_K) \cdot p(\text{open})$$
$$\gamma = 0.0236 \cdot [K^+]_o^{0.24}$$
$$p(\text{open}) = \frac{0.8}{1 + \left(\frac{[ATP]_i}{0.1} \right)^2}$$

Equation

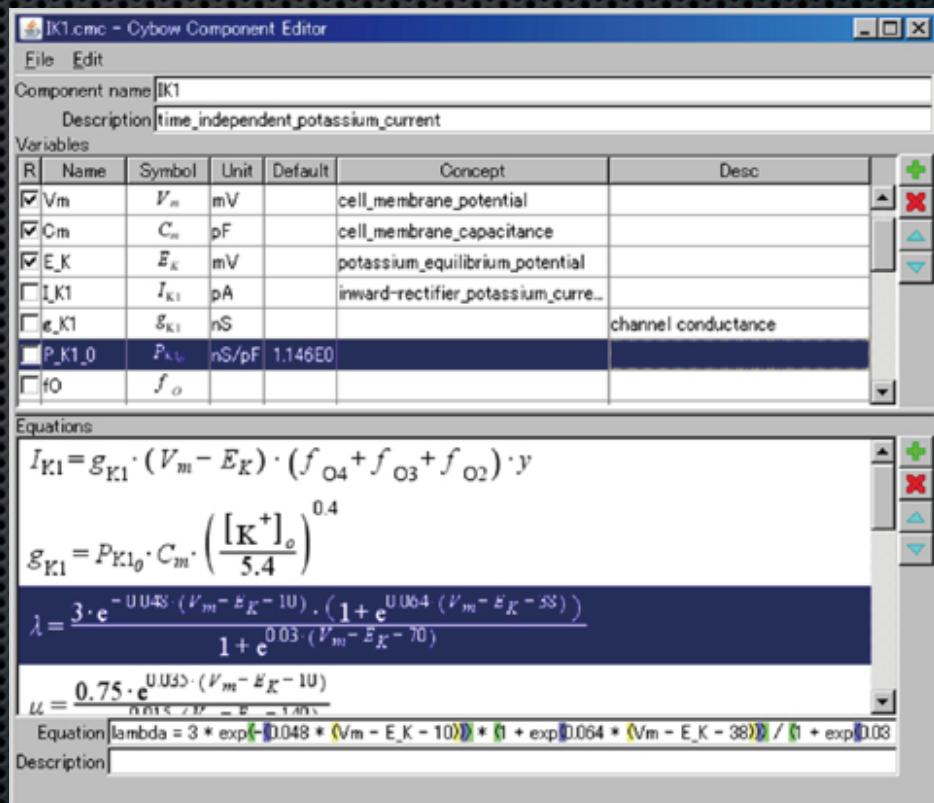
Description

Compose model



Component Editor

- Graphical rendering of math equations
- Simple one-line text input of a equation
- Graphical symbols for variables
- Annotation to model variables
- HTML document output
- Exportable equations to MS Office 2007

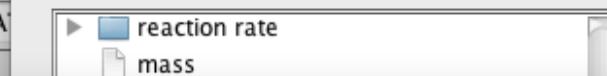


IKATP.cmc - Cybow Component Editor

Component Name K

Select Concept

Description A



potassium equilibrium potential
is-a-subclass-of. equilibrium potential

Variable

R	Name
<input type="checkbox"/>	N
<input type="checkbox"/>	gamma
<input checked="" type="checkbox"/>	Vm
<input checked="" type="checkbox"/>	EK
<input checked="" type="checkbox"/>	K_o
<input type="checkbox"/>	tau_P
<input type="checkbox"/>	I_KAT

Edit Concept

Concept: calcium binding rate of calmodulin

is-a: reaction rate constant

of-reaction

Concept: calcium binding to calmodulin

is-a: chemical reaction

Cancel

Reset

OK

$$p(\text{open}) = -\frac{1}{1 + \exp(-\frac{Vm - EK}{K_o})}$$

Equation p_o

- calcium equilibrium potential
- cell membrane potential
- Elastic Modulus
- capacitance
- amount of substance
- flux
- concentration

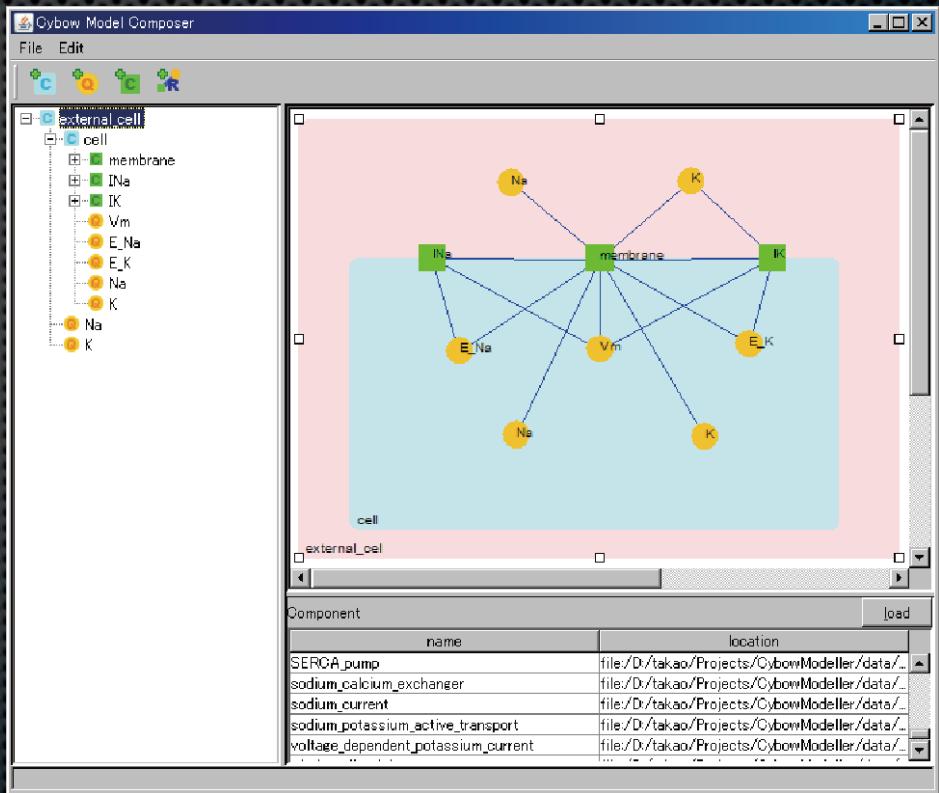
Cancel

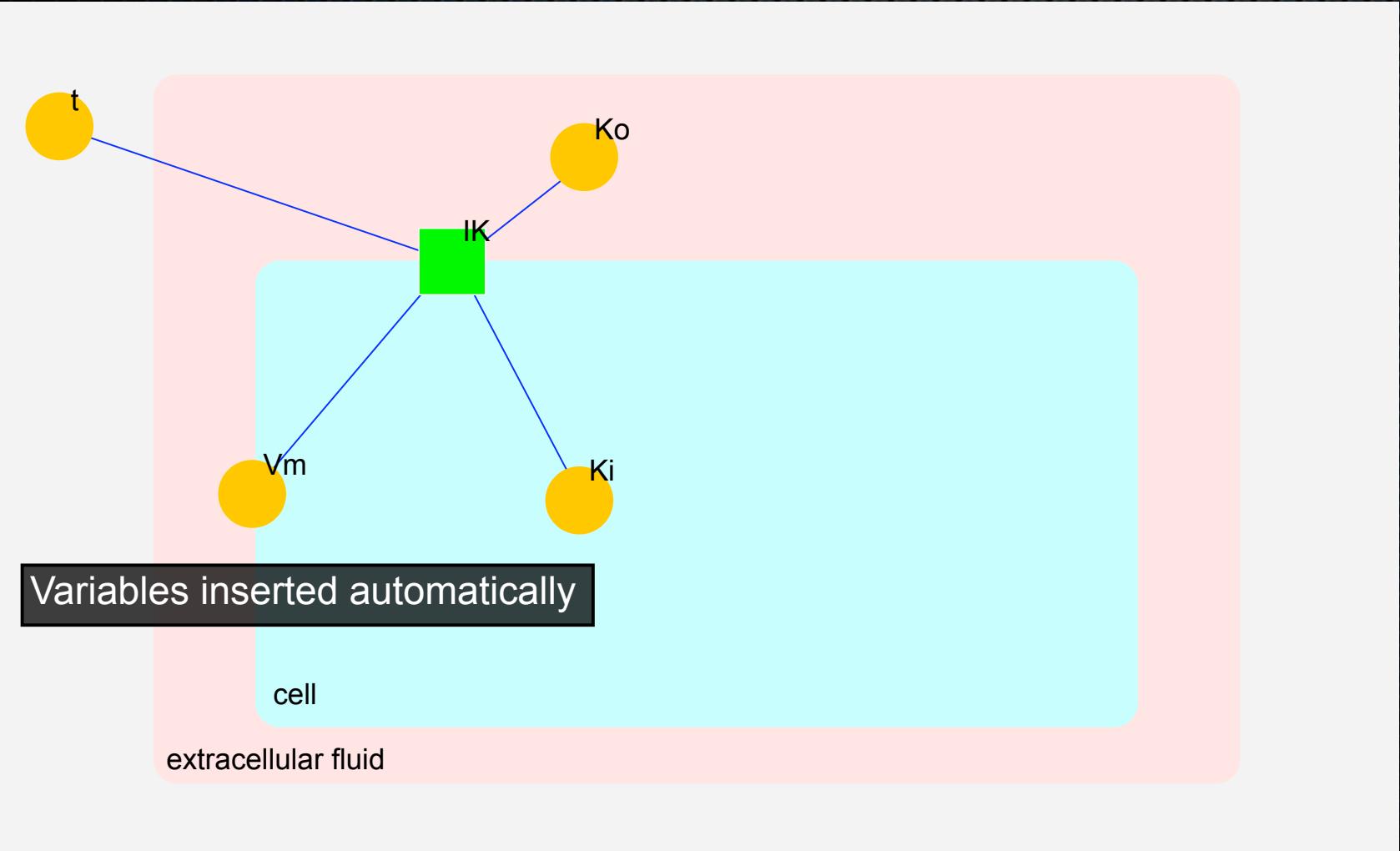
Define

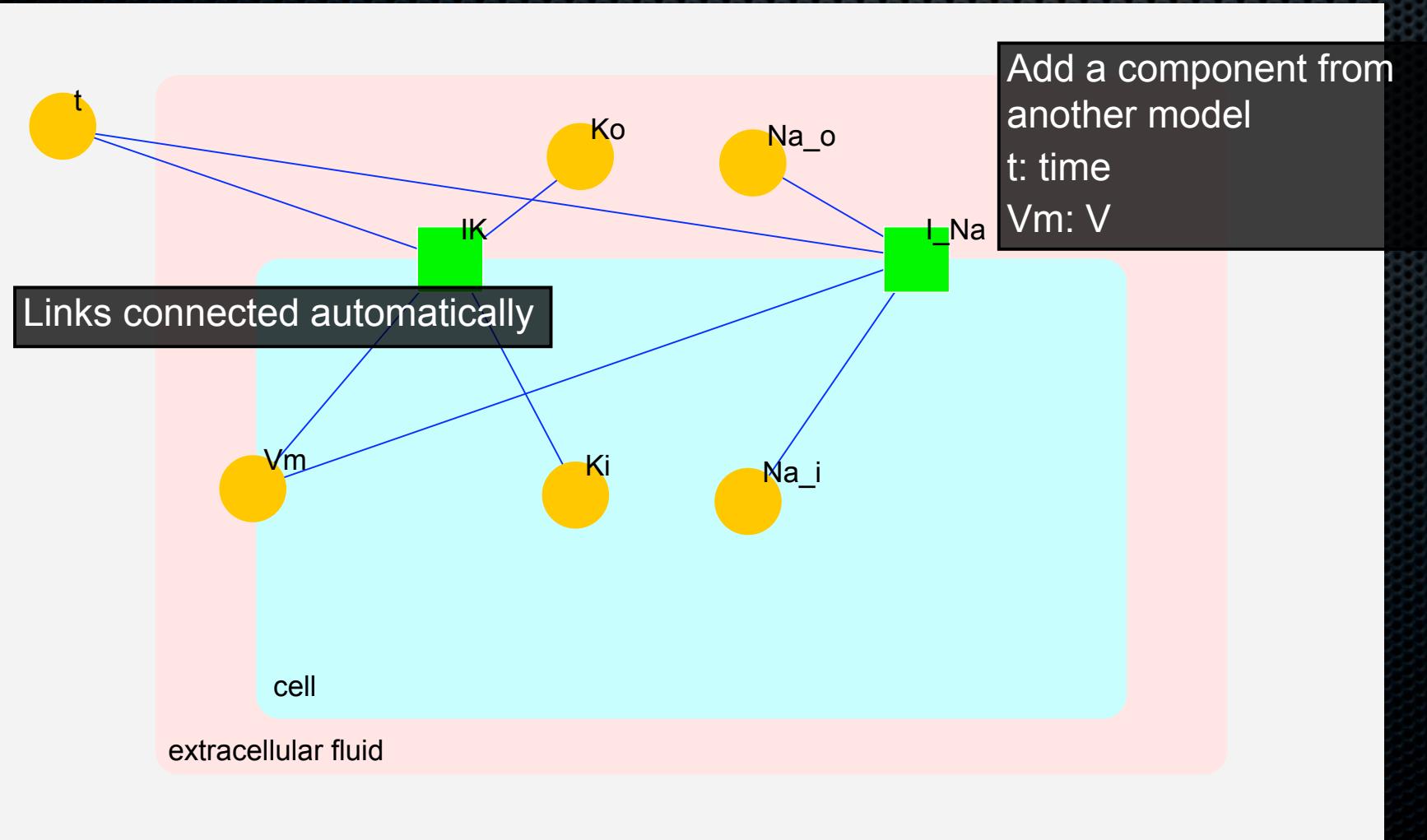
Select

Model Composer

- Graphical editing of model composition
- components & quantities on hierarchical compartments
- Automatic assistance by using annotations







Compatibility with CellML

- Original XML formats for components & compositions
 - Bindings are automatically generated from XSDs
- Translator from/to CellML in XSLT available
 - from CellML 1.0/1.1
 - import not supported
 - to CellML 1.1
 - variable symbols & annotations lost
 - not fully compatible yet

Code Generator

- C++ source codes, which uses LAPACK & CVODE
- Supports for semi-explicit DAE
 - doi: [10.1109/IEMBS.2009.5335041](https://doi.org/10.1109/IEMBS.2009.5335041)
- Not opened yet: prototype implementation

Summary

- Cybow Modeller: a software suite for cell modelling
 - Component Editor
 - Model Composer
 - CellML Translator
 - C++ Code Generator (under development)

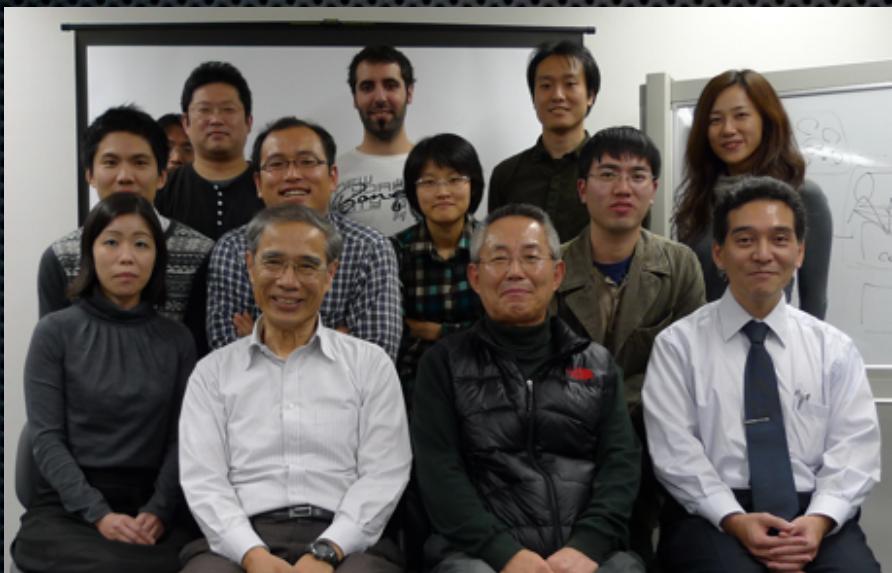
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Thank you!