RuleBuilder 1.0

Getting Started Guide

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Introduction

RuleBuilder enables you to define, generate, and simulate graphical rule-based models of biochemical systems. Objects and rules created in RuleBuilder are passed to the BioNetGen program in the form of a BioNetGen Language (BNGL) file in order to generate and simulate the reaction network. RuleBuilder enables you to control both the network generation and simulation steps and to view the generated network and plot the simulation results. This guide takes you through all of the basic steps involved in using RuleBuilder.

RuleBuilder Layout



Adding Containers and Components



Renaming Components and Containers



Resizing Containers



Creating Molecule Types

Molecules used in a model have to be defined and registered as a "Molecule Type" before they can be used in reaction rules and species.



Setting Allowed Component States

Components may take on different states to indicate conformation or covalent modification, such as phosphorylation.

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Allowed States for b	
A Default: Add Allowed State Add	Remove Done Component State dialog. Add an allowed state by typing in the Add Allowed State box and clicking Add. Components don't need to have any allowed states. Exit the dialog by clicking Done.
Object Manipulation Mode	•

Identifying Valid and Invalid Molecules



Copying Objects with the Selection Box

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Copying Objects with the Selection Box



Creating a Reaction Rule

Reaction rules are created by arranging containers and operators to construct a formula for the reaction.



Creating a Reaction Rule



Defining Products



Defining Products



Creating the Rule



Make Rule Dialog



Reaction Rules Window



Defining Seed Species

The network is defined by applying the reaction rules to a set of seed species.



Species Dialog Box

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	Species Properties Species Name Label: Species 1 Initial Concentration Parameter Name: B0 A dialog box appears for setting the name and initial concentration of the species.
A	Concentration: 1 Done Cancel
Object Manipulation Mode	

Seed Species Window



Defining Observables

Observables are concentration sums over species with particular properties and correspond to model outputs, such as total phosphorylation of a protein.



Make Observables Dialog



Observables Window



Observables Window



Running the Model

Once Reaction Rules, Seed Species, and Observables (optional) have been defined, the model can be simulated by pressing Run BioNetGen button.

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BioNetGen Engine Settings

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	These s Settings	ettings can also be changed by selecting under the File menu.	
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The Log Window

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Plotting the Results



Plotting the Results



Plotting the Results

