

4	The Design of SBML	6
4.1	SBML doesn't enforce units on math entities—are units for these entities not defined?	6
4.2	Why are reaction rates in "substance/time" units instead of "substance/volume/time" units?	7
4.3	How should models without compartments be encoded?	7
4.4	Why isn't there a default compartment with unit volume?	

1 SBML Introduction and Background

1.1 What is SBML?

The Systems Biology Markup Language (SBML) is a machine-readable format for describing qualitative and quantitative models of biochemical networks. It can also be used to express the interactions of biochemical networks with other phenomena. By a “biochemical network”, we mean a system consisting of biochemical

2.3 Are there large groups using SBML?

7.2 Who were the original SBML editors?

Michael Hucka, Herbert Sauro and Andrew Finney wrote the original 366(ed43(p)1(rop)-27(os)-1(a)1(c)27(k)in)-342(th)1(e