

JSBML: a flexible Java library for working with SBML

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ABSTRACT

Summary: The specifications of the Systems Biology Markup Language (SBML) define a standard for storing and exchanging biochemical models in XML-formatted text files. To perform higher-level operations on these models, e.g., numerical simulation or visual representation, an appropriate mapping to in-memory objects is required. To this end, the JSBML library has been developed. In contrast to earlier approaches, JSBML has been especially designed for the Java™ programming language and can therefore be used on all platforms, for which a Java Runtime Environment is available. In this way, programs based on JSBML can, for instance, be easily released as Java™ Web Start applications. JSBML's internal data structures have been developed entirely *de novo* based on the definitions in the SBML specifications. However, it strives to achieve the highest possible degree of compatibility to the existing library libSBML. JSBML supports all SBML levels and versions that are available today. In addition, JSBML provides modules that facilitate the development of plugins of applications such as CellDesigner or ease the migration from a libSBML backend.

Availability: Source code, binaries, and documentation of JSBML can be downloaded under the terms of LGPL 2.1 at <http://sbml.org/Software/JSBML>.

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Supplementary information: Supplementary data are available at Bioinformatics online.

1 INTRODUCTION

The XML-based Systems Biology Markup Language (SBML, Hucka *et al.* 2003) is the *de facto* standard file format for the storage and exchange of biochemical models, supported by more than 200 software packages to date (Feb. 2010). Much of this success is due to its clearly defined specifications and the availability of libSBML (Bornstein *et al.*, 2008), a portable, robust, and easy-to-use library.

LibSBML provides many methods for manipulating and validating SBML files through its Application Programming Interface (API). Originally written in C and C++, libSBML also provides automatically generated language bindings for Java™. However, the platform independence of Java is compromised in

libSBML due to the fact that the language binding is a wrapper around the C/C++ core. Therefore, many software developers experience difficulties in the deployment of portable libSBML-based Java applications. Because of this reason, among others, several groups in the SBML community started to develop their own native library for SBML. To avoid duplication of work, some of these groups have mounted an open-source effort to develop a pure Java library for SBML. Here we present the JSBML project, whose products are freely available at the web site <http://sbml.org/Software/JSBML>.

The aim of the project is to provide an SBML library that maps all SBML elements to a flexible and extended type hierarchy. Where possible, JSBML strives to attain 100 % compatibility with libSBML's Java API, to ease a switch from one library to the other. At the moment, there are no plans to re-implement some of the more complex functions of libSBML, such as model consistency checking, SBML validation, and the conversion between different SBML levels and versions, since separate community efforts are expected to make them available to JSBML via web services.

2 A BRIEF OVERVIEW OF JSBML

The main achievement of the JSBML project is its extended type hierarchy that has been designed from scratch based on the SBML specifications, but with respect to the naming conventions of methods and classes in libSBML. For each element defined in at least one of the SBML levels or versions, JSBML provides a corresponding class reflecting all of its properties. SBML elements or attributes that are not part of the most recent specifications (removed or obsolete) are marked as deprecated. JSBML defines a superclass or an interface for elements sharing common properties. For instance, the interface `NamedSBase` does not correspond to a data type in one of the SBML specifications directly, but serves as the superclass of all those instances of `SBase` that can be addressed by an identifier and a name. Similarly, all classes that may contain a mathematical expression implement the interface `MathContainer`. A full overview of this type hierarchy can be found in the supplementary file. JSBML also supports notes in XHTML form and annotations, including MIRIAM (Le Novère *et al.*, 2005) and SBO (Le Novère *et al.*, 2006). When building JSBML, the latest SBO file can directly be downloaded and parsed (Holland *et al.*, 2008). The `Model` class provides several `find*` methods to query for SBML elements. Filters enable to search lists for elements with certain properties. All `ListOf*` elements implement the Java's `List` interface, making it possible to iterate over its elements and to use generic types. Fig. 1 demonstrates how the

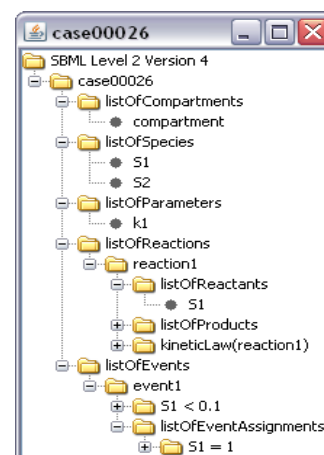
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```

1 import javax.swing.*;
2 import org.sbml.jsbml.*;
3
4 /** Displays the content of an SBML file in a {@link JTree} */
5 public class JSBMLvisualizer extends JFrame {
6
7     /** @param document The sbml root node of an SBML file */
8     public JSBMLvisualizer(SBMLDocument document) {
9         super(document.getModel().getId());
10        getContentPane().add(new JScrollPane(new JTree(document)));
11        pack();
12        setVisible(true);
13    }
14
15    /** @param args Expects a valid path to an SBML file. */
16    public static void main(String[] args) throws Exception {
17        new JSBMLvisualizer((new SBMLReader()).readSBML(args[0]));
18    }
19 }

```

(a) The SBML parser in JSBML considers the hierarchical data structure of SBML.



(b) Example for SBML test case 26

Fig. 1: Using JSBML for reading and visualizing an SBML file. In JSBML, the type `SBase` extends the Java interfaces `Serializable` for saving JSBML objects in binary form or sending them via a network connection, `Cloneable` to create deep object copies, and `TreeNode`. The latter one allows users to perform any kind of recursive operations on any instance of `SBase` such as visualizing it in a `JTree` (see 1b).

hierarchically structured content of an SBML file can be easily visualized in form of a tree. Special parsers read mathematical expressions from infix formulas or MathML into abstract syntax trees, which are the only internal representation of formulas in JSBML. These trees can directly be written in MathML, formula strings, or \LaTeX code. Although JSBML does not implement a full-featured consistency check for SBML models, some exceptions are thrown to prevent users to create invalid content. An overdetermination check for the model has been implemented based on the algorithm of Hopcroft and Karp (1973), which also identifies the variables in algebraic rules. Furthermore, JSBML can automatically derive the unit of a mathematical expression. Whenever a property of some `SBase` is altered, an `SBaseChangeEvent` is fired that notifies dedicated listeners. In this way, a graphical user interfaces could automatically react when any change happened to the model. With the help of some modules, JSBML capabilities can be extended and JSBML could be used as a communication layer between CellDesigner (Funahashi et al., 2003) or libSBML and any other applications. In this way, JSBML facilitates turning an existing application into a plug-in for CellDesigner. Currently, JSBML supports all constructs for SBML up to the latest Level 3 Version 1 Core specification.

3 IMPLEMENTATION

JSBML is entirely written in the Java™ version 1.5 and does not require the installation of any other software besides a Java Virtual Machine. JSBML is distributed in form of various JAR files (including or excluding required third-party libraries) and source code. In addition, a convenient build file with several options allows users to easily create customized JAR files. Being distributed under the terms of the Lesser GNU Public License (LGPL), the JSBML library can freely be used even in proprietary software.

4 CONCLUSION

JSBML is a young, ongoing software project that provides comprehensive and entirely Java-based data structures to read, write, and manipulate SBML files. Its layered architecture allows for the creation of Java web start applications and CellDesigner

plug-ins based on stand-alone programs with very little effort. One program, SBMLsqueezer (Dräger et al., 2008), has already been re-implemented and released under version 1.3 using JSBML. A simulator, which is benchmarked on the SBML test suite will be available soon, and many other projects are planned.

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Conflict of Interest: none declared.

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