# libsbml Developer's Manual

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## 1 Quickstart

LIBSBML depends on Apache's Xerces-C++ XML library for low-level XML tokenizing and Unicode support. Xerces is supported on Unix (Linux), Windows and MacOS X. Many popular Linux systems provide the Xerces library either as part of their standard distribution or as an optional RPM or Debian package. Apache provides a Windows binary distribution which includes both a DLL and a LIB file. For more information, see: http://xml.apache.org/xerces-c/. The instructions below assume Apache is already installed on your system.

#### 1.1 Linux, Cygwin or MacOS X

At the Unix, Cygwin or OS X command prompt, untar the distribution, cd into the directory created (e.g., libsbml-1.0.1/), and type:

```
% ./configure
% make
% make install
```

To compile programs that use libsbml (e.g., see Section 5.1) with GCC:

```
% gcc -o myapp.c myapp.c -lsbml
```

#### 1.2 Windows

Unzip the distribution and open the resulting folder (e.g. libsbml-1.0.1). There are debug (libsbmld) and release (libsbml) versions of LIBSBML, with .dll and .lib files for both versions in the Win32 subdirectory. LIBSBML header files are located in src/sbml.

Visual C++ projects should link with libsbml.lib or libsbmld.lib and generate code for the Multithreaded DLL or Debug Multithreaded DLL version of the VC++ runtime, respectively.

## 2 Introduction

This manual describes LIBSBML, a C application programming interface (API) library for reading, writing and manipulating the Systems Biology Markup Language (SBML; Hucka et al., 2001, 2003; ?). Currently, the library supports SBML Level 1 Version 1 and Version 2, and nearly all of SBML Level 2 Version 1. (The still-unimplemented parts of Level 2 are: support for RDF, and support for MathML's semantics, annotation and annotation-xml elements. These will be implemented in the near future.) For more information about SBML, please see the references or visit http://www.sbml.org/ on the Internet.

Since the library provides a C API, familiarity with the C programming language is assumed. Some parts of the library were written in C++, however, experience with C++ is not required; its use is "hidden" behind C functions. A companion document (Bornstein, 2003) provides a detailed reference manual for the API.

LIBSBML is entirely open-source and all specifications and source code are freely and publicly available. For more information about SBML, please see the references section or visit http://www.sbml.org/.

Some of the features of LIBSBML include:

#### • Small Memory Footprint

The parser is event-based (SAX2) and loads SBML data into C structures that mirror the classes in the SBML specification. No intermediate DOM is used, which greatly reduces runtime memory usage.

#### • Fast Runtime

The Gepasi generated 100 Yeast file (2Mb; 2000 reactions http://www.gepasi.org/gep3sbml.html) loads in 1.18s on a 1 GHz AMD Athlon XP and uses 1.4Mb of memory.

#### • Portable

The C source code is pure ANSI for maximum portability. It produces no errors or warnings when compiled with either GCC or Microsoft Visual C++. GCC compilation flags are: -ansi -pedantic-errors -Wall, i.e., ANSI violations are compilation errors instead of warnings and all warnings are reported.

The build system uses the GNU Autotools (Autoconf, Automake, and Libtool) to build shared and static libraries on a variety of Unix-based platforms.

Microsoft Windows is supported either through the Cygwin environment or as a native Win32 Dynamic Link Library (DLL). A Microsoft Visual C++ 6.0 project file is included in the source distribution and a precompiled Windows distribution is available.

#### • Full SBML Support

The parser supports both spellings of species and annotation (i.e., with and without the last s) for any level and version. When writing SBML documents, the appropriate specification-sanctioned spelling is used.

The full-text (including any namespace declarations) of <notes> and <annotation> elements may be retrieved from any SBML object. For compatibility with some technically incorrect but popular SBML documents, the parser recognizes and stores notes and annotations defined for the top-level <sbml> element (though a warning is logged).

#### • Full XML Schema Validation

The library uses the Apache Xerces-C++ XML library, which supports full XML Schema validation. All XML and Schema warning, error and fatal error messages are logged with line and column number information and may be retrieved and manipulated pragmatically. The Schema file to use for validation is configurable.

#### • Full Unicode Support

SBML Documents are parsed and manipulated in the Unicode codepage for efficiency (this is Xerces-C++ native format); however, strings are transcoded to the local code page for the SBML C structures.

#### • Well Tested

The entire library was written using the test-first approach popularized by Kent Beck and eXtreme Programming, where it's one of the 12 principles.

Currently there are 3014 individual assertions in 583 functional unit tests. Four test cases are responsible for reading entire SBML files (three are examples from the L1 document) into memory and verifying every field of the resulting structures.

#### • Conscientious Use of Memory

All memory for and contained in SBML structures is C memory, so realloc() and free() may be used. This is an important distinction when working with mixed C and C++ code. C's malloc() and free() are not compatible with C++'s new and delete operators.

A custom memory trace facility can be used to track all memory allocated and freed in both the library and all test suites. This facility must be enabled at build time with ./configure --enable-memory-tracing. For performance reasons memory tracing should be turned off in production environments. Currently all unit tests produce 1782 memory allocations and matched frees with no leaks.

#### 3 Installation

LIBSBML depends on Apache's Xerces-C++ XML library for low-level XML tokenizing and Unicode support. Xerces is supported on Unix (Linux), Windows and MacOS X. Many popular Linux systems provide the Xerces library either as part of their standard distribution or as an optional RPM or Debian package. Apache provides a Windows binary distribution which includes both a DLL and a LIB file.

For more information, see:

```
http://xml.apache.org/xerces-c/
```

A good way to determine whether or not Xerces-C is installed is to run the build script (see below); it will halt if it cannot find the Xerces-C library.

LIBSBML is designed to be extremely portable. It is written in 100% pure ANSI-C and the build system uses the GNU Autotools (Autoconf, Automake and Libtool). In most cases, building should be as easy unpacking the sources and running:

```
% ./configure
% make
% make check (optional)
% make install
```

However, if Xerces-C is installed in a non-standard place (e.g., a home directory), configure will not be able to detect it. In this case, configure needs to be told explicitly where to find the library. Set the CPPFLAGS and LDFLAGS environment variables to include the custom directories containing Xerces-C header and library files. For example:

```
% export CPPFLAGS=-I/home/bornstei/software/xerces-c/2.2.0/include
% export LDFLAGS=-L/home/bornstei/software/xerces-c/2.2.0/lib
% ./configure
% # etc.
```

make install copies header files to /usr/local/include/sbml and (shared and static) library files to /usr/local/lib. To specify a different install location use:

```
% ./configure --prefix=/my/favorite/path
```

make check is optional and will build and run an extensive suite of unit tests to verify all facets of the library. These tests are meant primarily for developers of LIBSBML and running them is not required for the library to function properly.

To run the unit tests a second library is required, libcheck. Check is a very lightweight C unit test framework based on the xUnit framework popularized by Kent Beck and eXtrememe Programming. Check is quite small and once installed, it consists of only two files: libcheck.a and check.h. To download Check, visit:

```
http://check.sf.net/
```

Note: Debian users can find Check as a standard add-on package (.deb).

All tests should pass with no failures or errors. If for some reason this is not the case on your system, please submit a bug report at http://www.sourceforge.net/projects/sbml.

#### 3.0.1 Memory Tracing

In addition to the unit tests, a custom memory tracing facility is available. It is disabled by default and must be enabled explicitly at build time, either as an argument to configure:

```
\% ./configure --enable-memory-tracing
```

or, in your own projects, by defining the C preprocessor symbol TRACE\_MEMORY:

```
#define TRACE_MEMORY
```

With memory tracing turned on, every piece of memory in both the library and all test suites is tracked. At the end of the run statistics are printed on total memory allocations, deallocations and leaks.

The memory statistics for the test suites should report zero leaks. If for some reason this is not the case, please submit a but report at http://www.sourceforge.net/projects/sbml.

For performance reasons, memory tracing should be disabled in production environments:

% ./configure --disable-memory-tracing

#### 4 SBML Classes in C

The SBML specification, with its UML diagrams, is suggestive of an object-oriented (OO) design. An API which attempts to represent SBML would do well to use an object-oriented programming (OOP) style to lower the inevitable impedance mismatch between specification and implementation. Unfortunately, the C programming language was not designed with OOP in mind and therefore does not support many of its features. It is possible, however, to construct a minimal object-like system in C with few, if any, drawbacks. For these reasons, the LIBSBML API mimics an object-oriented programming style.

The particular OOP-like style used by LIBSBML is not revolutionary. In fact, it is quite common and comprised of only a few simple stylistic conventions:

- 1. SBML classes are represented as C structs with a typedef shorthand. The shorthand form is derived by appending \_t to the name of the SBML class, e.g., Model becomes Model\_t.
- 2. C "objects" are nothing more than pointers to specific structs in memory. These pointers, instead of the structs themselves, are passed to and returned from "methods" (functions).
- 3. Functions meant to represent methods of (or messages to) an object are named beginning with the SBML class, followed by an underscore and ending in the method name. The functions take the object (pointer to struct) receiving the method as their first argument. For example, the function prototype for the addCompartment() method of a Model is:

```
void Model_addCompartment(Model_t *m, Compartment_t *c);
```

4. Constructor and destructor names are similar to method names, but end in \_create() and \_free(), respectively.

Every SBML class defined in the specification has a corresponding C class (see Table 1). The two SBML enumeration types, UnitKind and RuleType are represented as C enumerations, but deviate slightly from the above rules (see Section 4.5 for more information). Finally, there is one class, SBMLDocument\_t, that exists in the LIBSBML API, but not strictly in SBML. It serves primarily as a top-level container for models and stores any warning or error messages encountered when an SBML document was read (see Section 5).

The methods for SBML classes are declared in header files that correspond to the class name (e.g., Model.h). To include all methods for all classes in one fell swoop, #include SBMLTypes.h.

#### 4.1 Primitive Types

The mapping from SBML primitive types to C is straightforward as an example will help illustrate. A Species has at least one attribute of every primitive type defined by SBML.

SBML Class	C Class (typedef struct)
SBase	SBase_t
Model	$\mathtt{Model}_{\mathtt{-}}t$
UnitDefinition	${\tt UnitDefinition\_t}$
Unit	${\tt Unit\_t}$
Compartment	Compartment_t
Parameter	Parameter_t
Species	Species_t
Reaction	Reaction_t
SpeciesReference	SpeciesReference_t
KineticLaw	KineticLaw_t
Rule	Rule_t
As signment Rule	${\tt AssignmentRule\_t}$
AlgebraicRule	${\tt AlgebraicRule\_t}$
${\bf Compartment Volume Rule}$	${\tt CompartmentVolumeRule\_t}$
ParameterRule	$ParameterRule_t$
${\bf Species Concentration Rule}$	${\tt SpeciesConcentrationRule\_t}$
SBML Enumeration	C Enumeration (typedef enum)
UnitKind	UnitKind_t
RuleType	$RuleType_t$

**Table 1:** SBML classes and enumerations and their corresponding C class. Italicized classes are abstract, which sets their C implementation slightly apart from the others. See Section 4.6 for more information.

```
typedef struct
                        Species
name: SName
                                                                  char
                                                                          *name:
compartment: SName
                                                                  char
                                                                          *compartment;
initialAmount: double
                                                                  double initialAmount;
units: SName {use="optional"}
                                                                  char
                                                                          *units;
                                                                  int
                                                                          boundaryCondition;
boundaryCondition: boolean {use="optional" default="false"}
                                                                  int
                                                                          charge;
charge: integer {use="optional"}
                                                                } Species_t;
```

Figure 1: Definition of SBML Species in UML (left) and the corresponding Species\_t C struct (right).

Figure 1 shows the UML definition for Species and the corresponding C struct side-by-side. The similarity between the two demonstrates the mapping rules for primitive types:

- 1. In all cases, the form of UML attribute names, including their capitalization, are preserved (e.g., initialAmount) when mapped to C struct fields.
- 2. SName maps to a standard C string (pointer to an array of characters terminated by a NULL (0) character; for e.g. char \*name). Note that SName syntax is not yet enforced in the API.
- 3. double and integer map to C double and int respectively.
- 4. boolean maps to C int, where zero represents false and non-zero represents true.

#### 4.2 Object Creation and Destruction

This section and subsequent ones focus on functions or methods to create, destroy and otherwise manipulate SBML C objects. Since all functions and methods follow the same naming convention, when discussing them generically, XXX will be used to stand for some class name and YYY some class attribute.

To instantiate (create) an object use either the XXX\_create() or XXX\_createWith() constructor. To destroy (free) an object use XXX\_free(). The constructors and destructors for Species are:

#### Species\_t \*Species\_create (void)

Creates a new Species and returns a pointer to it.

# Species\_t \*Species\_createWith (const char \*name, const char \*compartment, double initialAmount, const char \*units, int boundaryCondition, int charge)

Creates a new Species with the given name, compartment, initialAmount, units, boundaryCondition and charge and returns a pointer to it. This convenience function is functionally equivalent to:

```
Species_t *s = Species_create();
Species_setName(s, name); Species_setCompartment(s, compartment); ...;
```

#### void Species\_free (Species\_t \*s)

Frees the given Species.

The XXX\_createWith() constructors are a convenient way both to create SBML objects and initialize their attributes in a single operation. If XXX\_create() is used instead, only attributes with default values (as defined by the specification) will be set. All other attributes will default to zero in the case of integers and doubles, false in the case of booleans or a NULL pointer in the case of SNames strings.

When an SBML object is destroyed with XXX\_free(), all of its Strings are freed (see Section 4.3: Accessing Fields for more information) and all of its contained objects are freed (see Section 4.4: Lists for more information).

#### 4.3 Accessing Fields

#### 4.3.1 Getters

There are two ways to get the value of a field within a struct: i) direct reference and ii) via a method call. Direct reference should be preferred in almost all cases; it avoids the overhead of a function call and thus is significantly faster<sup>1</sup>. The getter methods are provided for languages that do not allow direct access to the underlying C structs. As an example of direct access, Figure 2 lists a function (which is not part of the API) to print the fields of a Species.

The getter methods follow the naming convention XXX\_getYYY(). The getters for Species are:

```
const char *Species_getName (const Species_t *s)
Returns the name of this Species.
```

```
const char *Species_getCompartment (const Species_t *s)
Returns the compartment of this Species.
```

```
double Species_getInitialAmount (const Species_t *s)
Returns the initialAmount of this Species.
```

<sup>&</sup>lt;sup>1</sup>The getters are *not* #define macros that are inlined at compilation time. They are true functions.

```
1
2
    * Prints Species to stream (good for debugging).
3
    */
   void
4
   myPrintSpecies (Species_t *s, FILE *stream)
5
6
                         = "(none)";
7
     char none[]
     char *compartment = (s->compartment == NULL) ? none : s->compartment;
8
                                              == NULL) ? none : s->units;
9
     char *units
                         = (s-)units
                         = (s->boundaryCondition == 0) ? "false" : "true";
10
     char *boundary
11
12
     fprintf(stream, "Species %s:\n", s->name);
13
     fprintf(stream, "
                                  {\tt Compartment: \ \%s\n", \ compartment}
14
                               Initial Amount: %f\n", s->initialAmount );
     fprintf(stream,
15
     fprintf(stream,
                                                                            );
16
                                         Units: %s\n", units
                          Boundary Condition: %s\n", boundary Charge: %d\n", s->charge
17
     fprintf(stream,
                                                                            );
18
     fprintf(stream,
                                                                            );
   }
19
```

Figure 2: Demonstrates direct access to the fields of an SBML object, in this case Species\_t.

```
const char *Species_getUnits (const Species_t *s)
Returns the units of this Species.
```

```
int Species_getBoundaryCondition (const Species_t *s)
Returns the boundaryCondition of this Species.
```

```
int Species_getCharge (const Species_t *s)
Returns the charge of this Species.
```

Notice the Species\_t passed to each getter is constant. The purpose of this constness is twofold: i) it reinforces the notion that a getter simply returns a value and does not modify the state of the passed-in object and ii) as a result, in certain contexts a compiler may be able to use this information to perform certain optimizations. Notice also, whenever a getter returns a string, it is constant (const char \*), i.e. it cannot be modified or freed. The reason for this is each struct tracks and owns all of its internal memory. To modify (or especially free) this memory without using one of the sanctioned access methods could be particularly disasterous (most likely resulting in a segmentation or general protection fault). Memory management issues are elaborated when discussing setters.

#### 4.3.2 Setters

A value is assigned to a field via a set method<sup>2</sup>. Requiring all assignments to take place through a setter allows LIBSBML to track (and the developer to query) the set or unset *state* of a field apart from its actual *value*. The need to make a distinction between state and value is critical and is discussed further in Section 4.3.3.

The setter methods follow the naming convention XXX\_setYYY(). The setters for Species are:

```
void Species_setName (Species_t *s, const char *sname)
Sets the name of this Species to a copy of sname.
```

 $<sup>^2</sup>$ Earlier versions of LIBSBML allowed primitive types to be set directly. Such direct access, however, made it impossible to distinguish between set and unset states of a field, since all possible values are valid (i.e. no sentinel value exists to indicate an unset state).

#### void Species\_setCompartment (Species\_t \*s, const char \*sname)

Sets the compartment of this Species to a copy of sname.

#### void Species\_setInitialAmount (Species\_t \*s, double value)

Sets the initialAmount of this Species to value and marks the field as set.

#### void Species\_setUnits (Species\_t \*s, const char \*sname)

Sets the units of this Species to a copy of sname.

#### void Species\_setBoundaryCondition (Species\_t \*s, int value)

Sets the boundary Condition of this Species to value (boolean) and marks the field as set.

#### void Species\_setCharge (Species\_t \*s, int value)

Sets the charge of this Species to value and marks the field as set.

In the case of strings, requiring setter methods also enables clean and simple memory semantics. The rule is: every SBML object is responsible for its own memory, including SName strings. Whenever a set method is called, the passed-in string is copied and stored. If the field being set previously contained a string, it is freed. When XXX\_free() is called, all strings are freed.

For example, to set the compartment of Species s to the string "cell":

```
Species_setCompartment(s, "cell");
```

The effect of passing a NULL pointer as the string argument is to free the previously stored string and mark the field as unset. The preferred method for doing this, however, is to use the XXX\_unsetYYY() class of methods (see Section 4.3.3).

There is nothing in C language specification or the compiler that prevents string fields (or any other field type for that matter) from being set directly, as in:

```
s->name = "s2";
```

But to do so would likely cause a memory leak if s->name was already assigned to another string. Using the setter methods is much safer than assigning the memory directly and enables the set versus unset state of the field to be tracked.

#### 4.3.3 Field States

For each optional field without a default value, LIBSBML tracks both its state and value. The state of a field indicates whether the field is set (contains a valid value) or unset (contains no value at all). As mentioned before, the distinction between a set and unset field is critical for both LIBSBML and applications that depend upon it to function correctly (in accordance with the SBML specifications). Take for example the case of outputting SBML for a Species. Species has an optional charge attribute, which means it need not ever be read-in (specified), written or manipulated. For this reason, LIBSBML must be able to determine the set or unset state of the charge field to either output or omit the field as appropriate.

The easiest way to determine whether a particular field in a structure is set or unset is to use the XXX\_isSetYYY() class of methods. For Species, the following are available:

#### int Species\_isSetName (const Species\_t \*s)

Return 1 if the name of this Species has been set, 0 otherwise. In SBML L1, a Species name is required and therefore **should always be set**. In L2, name is optional and as such may or may not be set.

#### int Species\_isSetCompartment (const Species\_t \*s)

Return 1 if the compartment of this Species has been set, 0 otherwise.

#### int Species\_isSetInitialAmount (const Species\_t \*s)

Returns 1 if the initialAmount of this Species has been set, 0 otherwise. In SBML L1, a Species initialAmount is required and therefore **should always be set**. In L2, initialAmount is optional and as such may or may not be set.

#### int Species\_isSetUnits (const Species\_t \*s)

Returns 1 if the units of this Species has been set, 0 otherwise.

#### int Species\_isSetCharge (const Species\_t \*s)

Returns 1 if the charge of this Species has been set, 0 otherwise.

Fields with default values do not have an isSetYYY(). If the value for such a field is never supplied by an SBML document or user, the default is used. Therefore, if an isSetYYY() method did exist, it would always return true (1).

Required fields, on the other hand, do have isSetYYY() methods. There are two points worth mentioning here. First, it is possible that a value for a required field is not given and a program may want to check for and handle this case (especially if the program is an SBML validator). Second, please be aware from SBML Level 1 to Level 2, some fields changed from required to optional. If this is the case for a particular field the method documentation will state it (as above). That said, this second point is more relevant when it comes to unsetting fields.

Just as fields may be set and their set state queried, they may also be unset. Unset methods are named (predictably) XXX\_unsetYYY(). The unset methods for Species are:

#### void Species\_unsetName (Species\_t \*s)

Unsets the name of this Species. This is equivalent to the following code: safe\_free(s->name); s->name = NULL;

In SBML L1, a Species name is required and therefore **should always be set**. In L2, name is optional and as such may or may not be set.

#### void Species\_unsetInitialAmount (Species\_t \*s)

Unsets the initialAmount of this Species. In SBML L1, a Species initialAmount is required and therefore **should always be set**. In L2, initialAmount is optional and as such may or may not be set.

#### void Species\_unsetUnits (Species\_t \*s)

Unsets the units of this Species. This is equivalent to: safe\_free(s->units); s->units = NULL;

```
void Species_unsetCharge (Species_t *s)
Unsets the charge of this Species.
```

Again, for the reason mentioned above, fields with default values do not have unsetYYY() methods. Similarly, required fields have unsetYYY() methods only if they are declared optional in at least one of SBML Level 1 and Level 2. Notice, for example, there is an isSetCompartment() method but no corresponding unsetCompartment() (a compartment is required for a Species in both SBML Level 1 and Level 2).

It is also possible to check the set status of a field directly. For SName strings this is simple, unset fields will contain a NULL pointer:

```
s->units != NULL /* Equivalent to Species_isSetUnits(s). */
```

For primitive types things are only slightly more complicated. A nested isSet struct of one bitfield per primitive type is used to keep track of set states. The value of a bitfield field is one if corresponding field is set and zero otherwise. The isSet struct was omitted from the definition of Species\_t in Figure 1 to simplify the explanation of getter methods. A more complete definition of Species\_t, with its isSet nested struct is shown in Figure 3.

So, to determine if a given species s has a charge:

```
s->isSet.charge == 1 /* Equivalent to Species_isSetCharge(s) */
```

```
typedef struct
1
2
3
     char
               *name;
4
     char
               *compartment;
5
     double
               initialAmount;
6
     char
               *units;
7
     int
               boundaryCondition;
8
     int
               charge;
9
10
     struct
11
12
        unsigned int initialAmount:1;
13
        unsigned int charge
14
     } isSet;
15
   } Species_t;
16
```

**Figure 3:** Definition of Species\_t C struct, including its nested isSet struct of bitfields, used to track the state of primitive types.

#### 4.4 Lists

Species contains only SNames and primitive types, but many SBML classes also contain lists of other objects. For example, a UnitDefinition contains a list of Units:

```
    UnitDefinition
    Unit

    name : SName unit : Unit[0..*]
    kind : UnitKind exponent : integer {use="optional" default="1"} scale : integer {use="optional" default="0"}
```

To help manage this containment relationship, three standard functions are provided by LIB-SBML: XXX\_addYYY(), XXX\_getYYY() and XXX\_getNumYYY(). For example, the methods for UnitDefinition are:

```
void UnitDefinition_addUnit (UnitDefinition_t *ud, Unit_t *u)
```

Adds the given Unit to this UnitDefinition.

```
Unit_t *UnitDefinition_getUnit (const UnitDefinition_t *ud, unsigned int n)
```

Returns the nth Unit of this UnitDefinition.

```
unsigned\ int\ UnitDefinition\_getNumUnits\ (const\ UnitDefinition\_t\ *ud)
```

Return the number of Units in this UnitDefinition.

Furthering the example, creating the UnitDefinition  $mmoll^-1s^1$  named "mmls", corresponding to the SBML:

could be accomplished with the following C:

```
UnitDefinition_t *ud = UnitDefinition_createWith("mmls");
UnitDefinition_addUnit(ud, Unit_createWith(UNIT_KIND_MOLE , 1, -3) );
UnitDefinition_addUnit(ud, Unit_createWith(UNIT_KIND_LITRE , -1, 0) );
UnitDefinition_addUnit(ud, Unit_createWith(UNIT_KIND_SECOND, -1, 0) );
```

In this case, UnitDefinition\_getNumUnits(ud) returns 3 and UnitDefinition\_getUnit(ud, 1) returns the second Unit<sup>3</sup>. Notice that items are numbered starting at zero.

Related to lists is a set of convenience methods for creating and adding SBML objects to a Model in a single operation. The rationale is that since a Model is the top-level container for all other SBML objects, programmers are likely to have handles to them. Another way to construct the above UnitDefinition, but this time inside a Model, is:

```
Model_t          *m = Model_createWith("MyModel");
UnitDefinition_t *ud = Model_createUnitDefinition(m);
UnitDefinition_setName(ud, "mmls");

Model_createUnit(m, Unit_createWith(UNIT_KIND_MOLE , 1, -3) );
Model_createUnit(m, Unit_createWith(UNIT_KIND_LITRE , -1, 0) );
Model_createUnit(m, Unit_createWith(UNIT_KIND_SECOND, -1, 0) );
```

Model\_createUnit() creates a new Unit inside the Model m and returns a pointer to it (in this case the result is discarded). The Unit is added to the last UnitDefinition created. One caveat to be aware of with these methods is the case where no intermediate container exists; e.g., if no UnitDefinition were created above. In that case, the call to Model\_createUnit() does nothing. More specifically, no Unit is created and (obviously) nothing is added to the model.

For more detailed information on Lists, see Appendix A.

 $<sup>^3(\</sup>mbox{The UNIT\_KIND\_XXX}$  enumerations are discussed later.

#### 4.5 Enumerations

SBML has two enumeration types, UnitKind and RuleType. These translate directly to C enums with a few support functions for equality testing and converting to and from strings. From UnitKind.h:

```
typedef enum
{
    UNIT_KIND_AMPERE
, UNIT_KIND_BECQUEREL
, UNIT_KIND_CANDELA
    /* Omitted for space */
, UNIT_KIND_WATT
, UNIT_KIND_WEBER
, UNIT_KIND_INVALID
} UnitKind_t;
```

#### int UnitKind\_equals (UnitKind\_t uk1, UnitKind\_t uk2)

Tests for logical equality between two UnitKinds. This function behaves exactly like C's == operator, except for the following two cases:

- UNIT\_KIND\_LITER == UNIT\_KIND\_LITRE
- UNIT\_KIND\_METER == UNIT\_KIND\_METRE

where C would yield false (since each of the above is a distinct enumeration value), UnitKind\_equals(...) yields true. Returns true (!0) if uk1 is logically equivalent to uk2, false (0) otherwise.

#### UnitKind\_t UnitKind\_forName (const char \*name)

Returns the UnitKind with the given name (case-insensitive).

#### const char \*UnitKind\_toString (UnitKind\_t uk)

Returns the name of the given UnitKind. The caller does not own the returned string and is therefore not allowed to modify it.

The last item in the enumeration, UNIT\_KIND\_INVALID, is used whenever, as the name implies, the UnitKind is invalid or unknown. The corresponding string representation is "(Invalid UnitKind)". When a Unit is created, its kind field is initialized to UNIT\_KIND\_INVALID. Also, UnitKind\_forName() will return UNIT\_KIND\_INVALID if the passed-in name does not match any known UnitKind

The same ideas apply to RuleType, except there is no need for RuleType\_equals(). See RuleType.h for more information.

#### 4.5.1 Note:

The internal UNIT\_KIND\_STRINGS table is sorted alphabetically and UnitKind\_t matches this sort order. Because of this, UnitKind\_forName() is able to perform a binary search to find a matching name, making its complexity O(log(n)). That is, UnitKind\_forName() is implemented efficiently.

#### 4.6 Abstract Classes

The SBML specification defines three classes that have no representation apart from subclasses that specialize (inherit from) them. In OOP parlance, these types are termed abstract. The abstract SBML classes are:

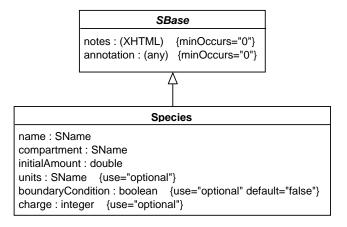
SBML Class	C Class (typedef struct)
SBase	SBase_t
Rule	${\tt Rule\_t}$
As signment Rule	${\tt AssignmentRule\_t}$

 Table 2: Abstract SBML classes their corresponding C class.

The conventions for abstract classes in the LIBSBML API are similar to that of other classes with a few modifications and additions.

Since abstract classes cannot be created or destroyed directly, they have no XXX\_create() or XXXX\_free() methods. Instead they have XXX\_init() and XXX\_clear() methods which subclasses use to initialize and free their memory, respectively. Users of the API do not need to worry about these functions.

Fields of the abstract class are #defined to a symbol in the class header file. This symbol is used in subclasses in the order of inheritance. For example, Species inherits from SBase:



SBase.h defines:

```
* As shown below, put SBASE_FIELDS as the *first* item of any struct
    * which "is a(n)" SBML object.
   #define SBASE_FIELDS
     SBMLTypeCode_t typecode; \
     char
                     *notes;
     char
                     *annotation
and recall Species_t from earlier:
   typedef struct
     SBASE_FIELDS;
     char
           *name;
     char
           *compartment;
     double initialAmount;
     char *units;
            boundaryCondition;
     int
```

```
int charge;
} Species_t;
```

The effect is that when the library source is compiled, the first three fields of Species are typecode, notes, and annotation. In fact, every class that inherits from SBase, i.e. all SBML classes, have these same first three fields. Accessing the notes or annotation field of a Species\_t \*s, or any other SBML object, is the same as for other fields. For example:

```
if (s->notes != NULL)
{
  printf("Notes for Species %s:\n", (s->name == NULL) ? "(null)" : s->name);
  printf("%s", s->notes);
}
```

Setting string fields requires special care to guard against memory leaks. The XXX\_setYYY() methods must be used. But, Species does not define either the notes or annotation fields and as such there are no Species\_setNotes() or Species\_setAnnotation() methods. Instead, SBase defines them:

```
void SBase_setNotes (SBase_t *sb, const char *notes)
```

Sets the notes field of the given SBML object to a copy of notes. If object already has notes, the existing string is freed before the new one is copied.

```
void SBase_setAnnotation (SBase_t *sb, const char *annotation)
```

Sets the annotation field of the given SBML object to a copy of annotations. If object already has an annotation, the existing string is freed before the new one is copied.

The first argument to these functions is, of course, an object of type SBase. Since Species inherits from SBase, i.e. Species is an SBase, it can be used as the first argument to these functions. A slight caveat is a cast is required. To set the notes field of Species s then:

```
SBase_setNotes( (SBase_t *) s, "My Favorite Species" );
```

The same applies to all other SBML objects.

Finally, each SBML class has a typecode which is initialized when an object is instantiated. The typecode is a simple C enumeration, defined in SBMLTypeCodes.h:

```
* An enumeration of SBML types to help identify SBML objects at runtime.
 * Abstract types do not have a typecode since they cannot be instantiated.
typedef enum
   SBML_COMPARTMENT
  , SBML_KINETIC_LAW
  , SBML_MODEL
  , SBML_PARAMETER
  , SBML_REACTION
  , SBML_SPECIES
  , SBML_SPECIES_REFERENCE
  , SBML_UNIT_DEFINITION
  , SBML_UNIT
  , SBML_ALGEBRAIC_RULE
  , SBML_SPECIES_CONCENTRATION_RULE
  , SBML_COMPARTMENT_VOLUME_RULE
   SBML_PARAMETER_RULE
} SBMLTypeCode_t;
```

The primary reason for the typecode is distinguish specific types of rules in a Model. A Model

contains a list of rules, but a Rule in SBML may be of one of four specific types: AlgebraicRule, SpeciesConcentrationRule, CompartmentVolumeRule and ParameterRule.

## 5 Reading SBML Files

SBML may be read from a file or an in memory string into an SBMLDocument. SBMLReader.h defines two basic read functions:

```
SBMLDocument_t *readSBML (const char *filename)
```

Reads the SBML document from the given file and returns a pointer to it.

#### SBMLDocument\_t \*readSBMLFromString (const char \*xml)

Reads the SBML document from the given XML string and returns a pointer to it. The XML string must be complete and legal XML document. Among other things, it must start with an XML processing instruction. For e.g.,:

```
<?xml version='1.0' encoding='UTF-8'?>
```

These functions return a pointer to an SBMLDocument:

```
/**
  * The SBMLDocument
  */
typedef struct
{
  unsigned int level;
  unsigned int version;
  List_t *error;
  List_t *fatal;
  List_t *warning;
  Model_t *model;
} SBMLDocument_t;
```

The level and version of the SBML document are stored in the first two fields. The last field is the SBML model itself. The three lists record warnings and errors encountered during the XML parse. Each warning or error is a ParseMessage (again from SBMLDocument.h):

```
/**
 * SBMLDocuments contain three Lists of ParseMessages, one for each class
 * of messages that could be triggered during an XML parse: Warnings,
 * Errors and Fatal Errors.
 *
 * Each ParseMessage contains the message itself and the line and column
 * numbers of the XML entity that triggered the message. If line or column
 * information is unavailable, -1 is used.
 */
typedef struct
{
    char *message;
    int line;
    int column;
} ParseMessage_t;
```

While its possible to access these lists directly, convenience functions are provided:

#### ParseMessage\_t \*SBMLDocument\_getWarning (SBMLDocument\_t \*d, unsigned int n)

Returns the nth warning encountered during the parse of this SBMLDocument or NULL if n > getNumWarnings() - 1.

#### ParseMessage\_t \*SBMLDocument\_getError (SBMLDocument\_t \*d, unsigned int n);

Returns the nth error encountered during the parse of this SBMLDocument or NULL if n > getNumErrors() - 1.

#### ParseMessage\_t \*SBMLDocument\_getFatal (SBMLDocument\_t \*d, unsigned int n)

Returns the nth fatal error encountered during the parse of this SBMLDocument or NULL if n > getNumErrors() - 1.

#### unsigned int SBMLDocument\_getNumWarnings (SBMLDocument\_t \*d)

Returns the number of warnings encountered during the parse of this SBMLDocument.

#### unsigned int SBMLDocument\_getNumErrors (SBMLDocument\_t \*d)

Returns the number of errors encountered during the parse of this SBMLDocument.

#### unsigned int SBMLDocument\_getNumFatals (SBMLDocument\_t \*d)

Returns the number of fatal errors encountered during the parse of this SBMLDocument.

#### void SBMLDocument\_printWarnings (SBMLDocument\_t \*d, FILE \*stream)

Prints all warnings encountered during the parse of this SBMLDocument to the given stream. If no warnings have occurred, i.e. SBMLDocument\_getNumWarnings(d) == 0, no output will be sent to stream. The format of the output is:

```
%d Warning(s):
Line %d, Col %d: %s
```

This is a convenience function to aid in debugging. For example: SBMLDocument\_printWarnings(d, stdout).

#### void SBMLDocument\_printErrors (SBMLDocument\_t \*d, FILE \*stream)

Prints all errors encountered during the parse of this SBMLDocument to the given stream. If no errors have occurred, that is, if SBMLDocument\_getNumErrors(d) == 0, no output will be sent to stream. The format of the output is:

```
%d Error(s):
Line %d, Col %d: %s
```

This is a convenience function to aid in debugging. For example: SBMLDocument\_printErrors(d, stdout).

#### void SBMLDocument\_printFatals (SBMLDocument\_t \*d, FILE \*stream)

Prints all fatals encountered during the parse of this SBMLDocument to the given stream. If no fatals have occurred, that is, if SBMLDocument\_getNumFatals(d) == 0, no output will be sent to stream. The format of the output is:

```
%d Fatal(s):
  Line %d, Col %d: %s
```

This is a convenience function to aid in debugging. For example: SBMLDocument\_printFatals(d, stdout).

#### 5.1 A Simple Example

The following example is included in the distribution as readSBML.c. It is compiled as part of the build process, but is not installed. The program takes a single command-line argument, the name of an SBML file, reads it into memory and reports some basic information about the file: warnings or errors (if any are reported) as well as the total read time (in milliseconds).

To run the example, go to the top-level directory where LIBSBML was unpacked and:

```
% cd src
% ./readSBML test-data/l1v1-branch.xml
% ./readSBML test-data/l1v1-minimal.xml
% ./readSBML test-data/l1v1-rules.xml
% ./readSBML test-data/l1v1-units.xml
% # etc...
```

The exact procedure for compile the example (and linking-in LIBSBML in general) varies from one platform and compiler to another. On Linux or Solaris with GCC the following should work:

```
% gcc -o readSBML readSBML.c -lsbml
```

The example is listed below:

```
#include <stdio.h>
 1
2
   #include <sys/time.h>
3
4
   #include "sbml/SBMLReader.h"
   #include "sbml/SBMLTypes.h"
5
6
7
8
    * Function Prototypes
   unsigned long getCurrentMillis (void);
10
                  reportErrors
                                     (SBMLDocument_t *d);
12
13
14
   main (int argc, char *argv[])
15
16
17
     SBMLDocument_t *d;
18
     Model_t *m;
19
20
     unsigned long start, stop;
21
22
23
     if (argc != 2)
24
       printf("usage: readSBML <filename>\n");
25
26
       return 1;
     }
27
```

```
28
29
     start = getCurrentMillis();
            = readSBML(argv[1]);
30
31
     stop = getCurrentMillis();
32
33
     m = d-> model;
34
     printf( "File: %s\n", argv[1]);
35
                       model name: %s\n",
36
     printf( "
                                            m->name );
     printf( "
                 unitDefinitions: %d\n",
                                            Model_getNumUnitDefinitions(m) );
37
     printf( "
                    compartments: %d\n",
                                            Model_getNumCompartments(m)
38
                                                                               );
                          species: %d\n",
     printf( "
39
                                             Model_getNumSpecies(m)
                                                                               );
                       parameters: %d\n",
  reactions: %d\n",
     printf( "
40
                                             Model_getNumParameters(m)
                                                                               );
     printf( "
                                             Model_getNumReactions(m)
41
                                                                               );
     printf( "
                            rules: %d\n",
                                             Model_getNumRules(m)
42
                                                                               );
     printf( "\n");
43
44
     printf( "Total Read Time (ms): %lu\n", stop - start);
45
46
47
     SBMLDocument_printWarnings(d, stdout);
48
     SBMLDocument_printErrors (d, stdout);
     SBMLDocument_printFatals (d, stdout);
49
50
51
     SBMLDocument_free(d);
52
     return 0;
53
   }
54
55
56
    * @return the number of milliseconds elapsed since the Epoch.
57
58
   unsigned long
59
   getCurrentMillis (void)
60
61
62
     struct timeval tv;
63
     unsigned long result = 0;
64
65
66
     if (gettimeofday(&tv, NULL) == 0)
67
        result = (tv.tv_sec * 1000) + (tv.tv_usec * .001);
68
69
70
71
     return result;
   }
72
```

#### 5.2 XML Schema Validation

To validate an SBML document against an SBML (XML) schema as it is being read-in requires creating an SBMLReader object and setting the appropriate schema filename and validation level. The functions for doing this are:

#### SBMLReader\_t \*SBMLReader\_create (void)

Creates a new SBMLReader and returns a pointer to it. By default schema validation is off (XML\_SCHEMA\_VALIDATION\_NONE) and schemaFilename is NULL.

#### void SBMLReader\_free (SBMLReader\_t \*sr)

Frees the given SBMLReader.

#### void SBMLReader\_setSchemaFilename (SBMLReader\_t \*sr, const char \*filename)

Sets the schema filename used by this SBMLReader. The filename should be either i) an absolute path or ii) relative to the directory contain the SBML file(s) to be read.

# $\label{lem:condition} {\tt Void SBMLReader\_setSchemaValidationLevel (SBMLReader\_t *sr, XMLSchemaValidation\_t level)}$

Sets the schema validation level used by this SBMLReader. The levels are:

- XML\_SCHEMA\_VALIDATION\_NONE (0) turns schema validation off.
- XML\_SCHEMA\_VALIDATION\_BASIC (1) validates an XML instance document against an XML Schema. Those who wish to perform schema checking on SBML documents should use this option.
- XML\_SCHEMA\_VALIDATION\_FULL (2) validates both the instance document itself and the XML Schema document. The XML Schema document is checked for violation of particle unique attribution constraints and particle derivation restrictions, which is both time-consuming and memory intensive.

Once an SBMLReader has been created, there two variants of the functions readSBML() and readSBMLFromString() become available. These variants can be thought of as methods of the SBMLReader class.

```
SBMLDocument_t *SBMLReader_readSBML (SBMLReader_t *sr, const char *filename)
```

Reads the SBML document from the given file and returns a pointer to it.

```
SBMLDocument\_t \ *SBMLReader\_readSBMLFromString \ (SBMLReader\_t \ *sr, \ const \ char \\ *xml)
```

Reads the SBML document from the given XML string and returns a pointer to it. The XML string must be complete and legal XML document. Among other things, it must start with an XML processing instruction. For e.g.,:

```
<?xml version='1.0' encoding='UTF-8'?>
```

Schema violations are reported in the SBMLDocument's list of ParseMessages.

# 6 Writing SBML Files

The SBMLWriter class is similiar to the SBMLReader class. The difference with an SBMLWriter is instead of selecting a schema filename and validation level, an output encoding is chosen instead. The supported output encodings are:

```
/**
  * The character encodings supported by SBMLWriter
  */
typedef enum
{
    CHARACTER_ENCODING_ASCII
, CHARACTER_ENCODING_UTF_8
, CHARACTER_ENCODING_UTF_16
, CHARACTER_ENCODING_ISO_8859_1
, CHARACTER_ENCODING_INVALID
} CharacterEncoding_t;
```

SBMLWriter.h defines the following functions:

#### SBMLWriter\_t \*SBMLWriter\_create (void)

Creates a new SBMLWriter and returns a pointer to it. By default the character encoding is ISO 8859-1 (CHARACTER\_ENCODING\_ISO\_8859\_1).

#### void SBMLWriter\_free (SBMLWriter\_t \*sw)

Frees the given SBMLWriter.

#### void SBMLWriter\_setEncoding (SBMLWriter\_t \*sw, CharacterEncoding\_t encoding)

Sets the character encoding for this SBMLWriter to the given CharacterEncoding type.

# int SBMLWriter\_writeSBML (SBMLWriter\_t \*sw, SBMLDocument\_t \*d, const char \*filename )

Writes the given SBML document to filename (with the settings provided by this SBML-Writer).

Returns 1 on success and 0 on failure (e.g., if filename could not be opened for writing or the SBMLWriter character encoding is invalid).

#### char \*SBMLWriter\_writeSBMLToString (SBMLWriter\_t \*sw, SBMLDocument\_t \*d)

Writes the given SBML document to an in-memory string (with the settings provided by this SBMLWriter) and returns a pointer to it. The string is owned by the caller and should be freed (with free()) when no longer needed.

Returns NULL on failure (e.g., if the SBMLWriter character encoding is invalid).

Finally, if the default character encoding of ISO 8859-1 is acceptable, there are two simpler equivalents to the above functions:

#### int writeSBML (SBMLDocument\_t \*d, const char \*filename)

Writes the given SBML document to filename with the settings provided by this SBML-Writer. This convenience function is functionally equivalent to:

```
SBMLWriter_writeSBML(SBMLWriter_create(), d, filename);
```

Returns 1 on success and 0 on failure (e.g., if filename could not be opened for writing or the SBMLWriter character encoding is invalid).

#### char \*writeSBMLToString (SBMLDocument\_t \*d)

Writes the given SBML document to an in-memory string (with the settings provided by this SBMLWriter) and returns a pointer to it. The string is owned by the caller and should be freed (with free()) when no longer needed. This convenience function is functionally equivalent to:

```
SBMLWriter_writeSBMLToString(SBMLWriter_create(), d);
```

Returns NULL on failure (e.g., if the SBMLWriter character encoding is invalid).

# 7 Handling of Mathematical Formulas and MathML

LIBSBML can read and write MathML 2.0 (W3C, 2000) content in SBML documents and data streams, as well as translate between MathML and the text-string formulas used in SBML Level 1. This section describes the MathML and mathematics handling capabilities of the library.

#### 7.1 Reading and Writing Formulas in Text-String Form

In SBML Level 1, mathematical formulas are expressed as text strings using a simple C-like syntax. In SBML Level 2, mathematical formulas are expressed in MathML syntax. The API interfaces to SBML files and data streams work with ASCII text strings in both cases; the difference between Level 1 and Level 2 mathematical expressions is entirely in the syntax of the strings.

Whether formulas in SBML are expressed in the C-like syntax of Level 1 or in the MathML syntax of Level 2, they are converted internally in LIBSBML into Abstract Syntax Trees (ASTs). The structure of ASTs is described in detail in Appendix B. When LIBSBML reads an SBML model, it converts the expressions into ASTs and stores the ASTs in the corresponding data structures that have mathematical formulas (such as in a KineticLaw). Thus, the KineticLaw\_getMath() method, for example, returns a pointer to the root of an AST corresponding to the formula stored there.

Many software packages provide users with the ability to express formulas for such things as reaction rate expressions, and these packages' interfaces often let users type in the formulas directly as strings. LIBSBML provides two high-level functions for working with mathematical expressions in the form of strings: SBML\_parseFormula and SBML\_formulaToString.

#### ASTNode\_t \*SBML\_parseFormula (const char \*formula)

Parses the given string as a mathematical formula in SBML Level 1 syntax form, and returns a representation of it as an Abstract Syntax Tree (AST). This function returns the root node of the AST. If the formula contains a syntax error, this function returns NULL instead.

#### char \*SBML\_formulaToString (ASTNode\_t \*tree)

Returns a text-string mathematical expression corresponding to the Abstract Syntax Tree given as the argument. The caller owns the memory allocated for the returned string and is responsible for freeing it when it is no longer needed.

Using these methods is exceedingly simple. The following is a code fragment that illustrates calling the parser function repeatedly with different formula strings, taking the ASTs returned each time and handing them back to the formula generator and comparing the strings to make sure they matched. (This is not something a real application would ever need to do, but it does simply illustrate the use of these two methods.)

```
const char *formulae[] =
{
    "1",
    "2.1",
    "2.1e+10",
    "foo",
    "1 + foo",
    "1 + 2 * 3",
    "(1 - 2) * 3",
    "1 + -2 / 3",
```

```
"1 + -2e-100 / 3",
  "1 - -foo / 3",
  "2 * foo^bar + 3.1",
  "foo()",
  "foo(1)",
  "foo(1, bar)",
  "foo(1, bar, 2<sup>-3</sup>)",
};
ASTNode_t *n;
char
         *s;
int
          i;
for (i = 0; i < *formulae[i]; i++)</pre>
 n = SBML_parseFormula( formulae[i] ); /* Convert string to AST */
 s = SBML_formulaToString(n);
                                           /* Convert AST back to string */
 fail_unless( !strcmp(s, formulae[i]), NULL );
 ASTNode_free(n);
 safe_free(s);
```

#### 7.2 Reading Formulas in MathML Form

There may arise situations in which an application needs to convert MathML directly into an AST. LIBSBML provides the utility function readMathMLFromString() for this purpose.

#### MathMLDocument\_t \*readMathMLFromString (const char \*xml)

Reads a string containing an XML MathML expression, constructs the corresponding Abstract Syntax Tree and returns a pointer to a MathMLDocument\_t holding the tree structure.

The object returned by readMathMLFromString() is a simple container for an AST. The class of this object, MathMLDocument, is not defined by the SBML language standard but is provided in LIBSBML as a utility class. MathMLDocument serves as a top-level container for XML documents containing only MathML; in some ways it mirrors the SBMLDocument class, which acts as a container for XML documents containing SBML. The definition of MathMLDocument\_t is as follows:

```
/**
  * The MathMLDocument
  */
typedef struct
{
  ASTNode_t *math;
} MathMLDocument_t;
```

The following are the functions defined for the MathMLDocument class:

# MathMLDocument\_t \*MathMLDocument\_create () Creates a MathMLDocument\_t object structure.

```
void MathMLDocument_free (MathMLDocument_t *d)
Frees the given MathMLDocument_t structure.
```

#### ASTNode\_t \*MathMLDocument\_getMath (const MathMLDocument\_t \*d)

Returns the Abstract Syntax Tree representation of the mathematical formula stored in this  ${\tt MathMLDocument\_t}$  structure.

Note that because the content passed to <code>readMathMLFromString()</code> is handed to an XML parser, the string given as argument must be a complete XML document. The following example illustrates the use of this function with a valid MathML input.

#### 7.3 Special Cases in Mathematical Formulas

#### **A** Lists

While List convenience methods (e.g., XXX\_getNumYYY()) are provided for every class, it is possible to access and manipulate each list directly. All lists are themselves objects of type List\_t. The full set of list methods are:

```
void List_add (List_t *list, void *item)
Adds item to this List.

void *List_get (List_t *list, unsigned int n)
```

```
Returns the nth item in this List. If n > List_size(list) - 1 returns NULL.

void *List_remove (List_t *list, unsigned int n)
```

```
Removes the nth item from this List and returns a pointer to it. If n > List_size(list) - 1 returns NULL.
```

```
unsigned int List_size (List_t *list)

Returns the number of elements in this List.
```

Since UnitDefinitions maintains a List of Units, the UnitDefinition example in Section 4.4 could also be written as:

```
UnitDefinition_t *ud = UnitDefinition_createWith("mmls");
List_add(ud->unit, Unit_createWith(UNIT_KIND_MOLE , 1, -3) );
List_add(ud->unit, Unit_createWith(UNIT_KIND_LITRE , -1, 0) );
List_add(ud->unit, Unit_createWith(UNIT_KIND_SECOND, -1, 0) );
```

However, this approach is not preferred. The best reason to use specific XXX\_getYYY() methods over the List API is that the former are typed to specific items, whereas List\_get() returns a void pointer that must be cast to a specific type. For example, compare:

```
Unit_t *u = UnitDefinition_getUnit(u, 1);
to:
Unit_t *u = (Unit_t *) List_get(ud->unit, 1);
```

Further, the first is more readable. The List API is mentioned i) for the sake of completeness and ii) currently the only way to remove an item from a list is to use the API directly:

```
List_remove(ud->unit, 2);
```

has no analog in UnitDefinition. "Remove" convenience methods can be added to the API. This feature was skipped because list item removal seemed like an uncommon operation.

# B Abstract Syntax Trees and ASTNode\_t

Abstract Syntax Trees (ASTs) in LIBSBML are a simple data structure for storing mathematical expressions. For many applications, the details of ASTs are irrelevant because the applications can use the text-string based translation functions described in Sections 7.1 and 7.2. However, other applications do need to read and manipulate ASTs directly. This section describes LIBSBML's AST in sufficient detail so software authors can write code to work with them.

An AST *node* is a recursive structure containing a pointer to the node's value (e.g., a number or a symbol) and a list of children nodes. The following is the definition of ASTNode\_t:

```
* An Abstract Syntax Tree (AST) node.
typedef struct
 ASTNodeType_t type;
 union
   char
          ch;
   char
          *name;
   long
         integer;
   double real;
  } value;
  union
   long denominator;
   long exponent;
  } extra;
 List_t *children;
} ASTNode_t;
```

For nodes containing character, string, integer, or simple floating point data, the value union in ASTNode\_t holds the value directly. The additional field extra is used to handle two special cases that arise from the need to handle MathML:

• A numerical value represented in MathML as a real number with an exponent is stored in an AST node by putting the mantissa part in the field value.real and the exponent part in extra.exponent. Thus, a MathML value such as "1.0e4" will be stored in an AST node with value.real equal to 1.0 and extra.exponent equal to 4.

Readers may note that some numbers could be stored in a C double data type even if they were expressed in e-notation in MathML. Nevertheless, LIBSBML records these numbers in e-notation in the AST node so that when an SBML model is read in and then written out again, the amount of change introduced to the SBML during the round-trip activity is minimized.

 Rational numbers are represented in an AST node using the field value.integer to record the numerator and extra.denominator to record the denominator.

The use of unions for value and extra is for space efficiency only; the quantities recorded in them are mutually exclusive and so there is no need to assign separate storage to each one in the AST data structure.

The children field of ASTNode\_t is a list of pointers to other ASTNode\_t objects. This list is empty for AST nodes that are leaf elements, such as numbers. For AST nodes that are actually roots of expression subtrees, the list of children points to the parsed objects that make up the rest of the expression.

The following is a simple example.

#### NEED EXAMPLE

LIBSBML defines a number of functions for working with ASTNode\_t objects; see the LIBSBML API Reference Manual for a detailed list of these.

### References

Bornstein, B. (2003). LIBSBML API reference manual. Available via the World Wide Web at http://www.sbml.org.

Hucka, M., Finney, A., Sauro, H. M., and Bolouri, H. (2001). Systems Biology Markup Language (SBML) Level 1: Structures and facilities for basic model definitions. Available via the World Wide Web at http://www.sbml.org.

Hucka, M., Finney, A., Sauro, H. M., Bolouri, H., Doyle, J. C., Kitano, H., Arkin, A. P., Bornstein, B. J., Bray, D., Cornish-Bowden, A., Cuellar, A. A., Dronov, S., Gilles, E. D., Ginkel, M., Gor, V., Goryanin, I. I., Hedley, W. J., Hodgman, T. C., Hofmeyr, J.-H., Hunter, P. J., Juty, N. S., Kasberger, J. L., Kremling, A., Kummer, U., Le Novère, N., Loew, L. M., Lucio, D., Mendes, P., Minch, E., Mjolsness, E. D., Nakayama, Y., Nelson, M. R., Nielsen, P. F., Sakurada, T., Schaff, J. C., Shapiro, B. E., Shimizu, T. S., Spence, H. D., Stelling, J., Takahashi, K., Tomita, M., Wagner, J., and Wang, J. (2003). The Systems Biology Markup Language (SBML): A medium for representation and exchange of biochemical network models. *Bioinformatics*, 19(4):524–531.

W3C (2000). W3C's math home page. Available via the World Wide Web at http://www.w3.org/Math/.