SBML L3 Package Development

Sarah Keating





EBI is an Outstation of the European Molecular Biology Laboratory.

Outline

- 1. Brief overview of what SBML is and why it was developed
- 2. Overview of SBML Level 3 modularisation
- 3. L3 package development
- 4. libSBML and L3 packages
- 5. Links and acknowledgements





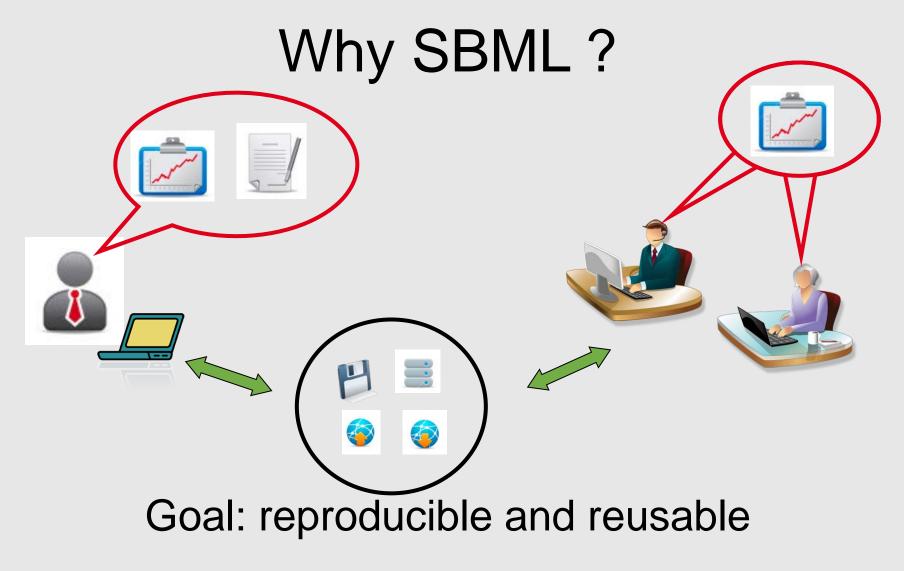
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models and simulations





What is SBML?

A machine-readable format for representing computational models in systems biology

Tool-neutral exchange language for software applications

Expressed in XML

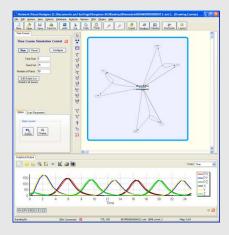
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What is SBML?

- Independent of modelling formalism
- Declares model not procedure







Outline

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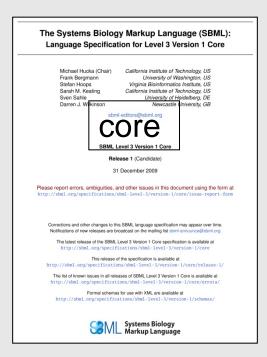
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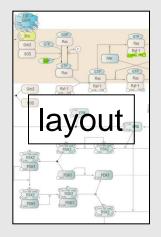


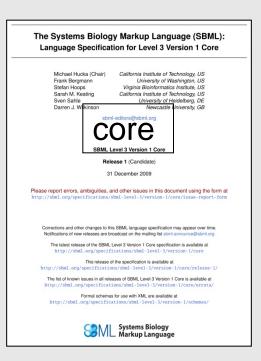






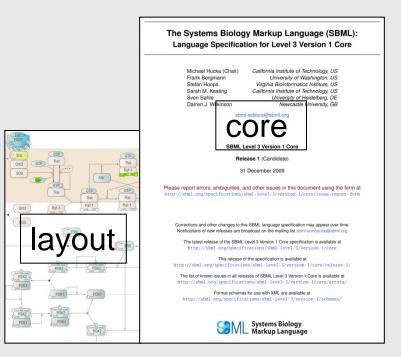






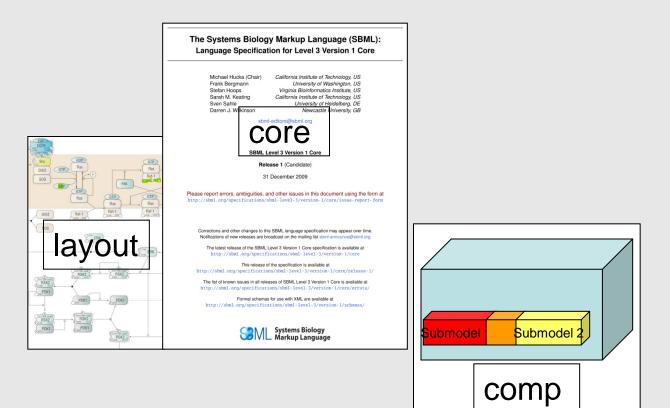






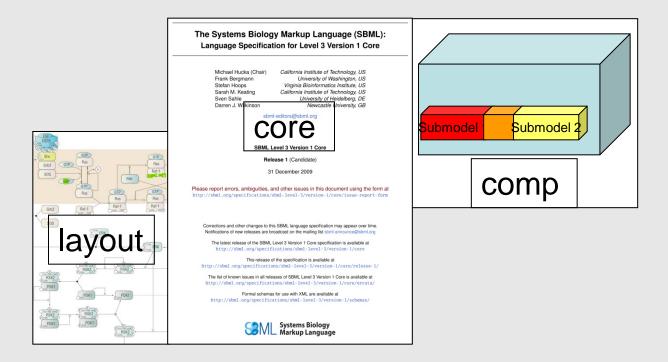






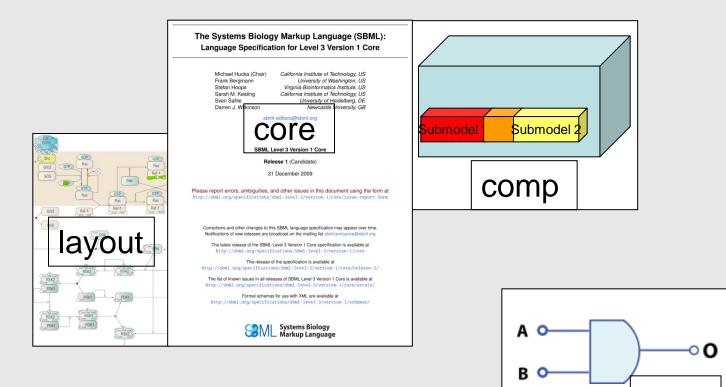








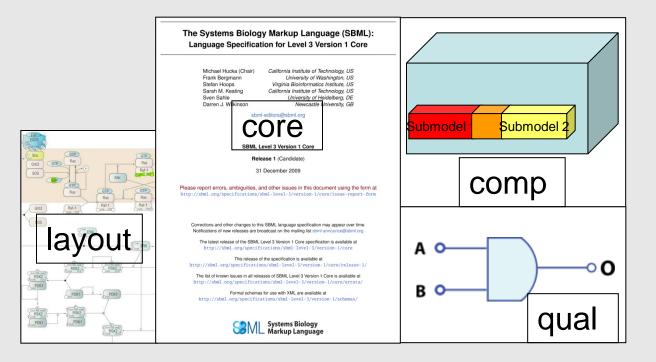






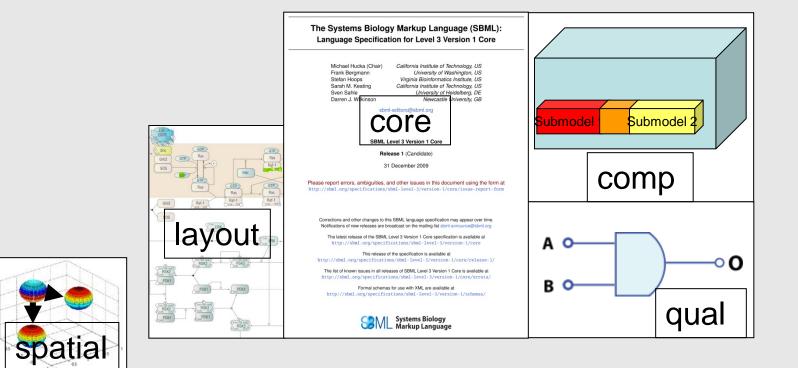


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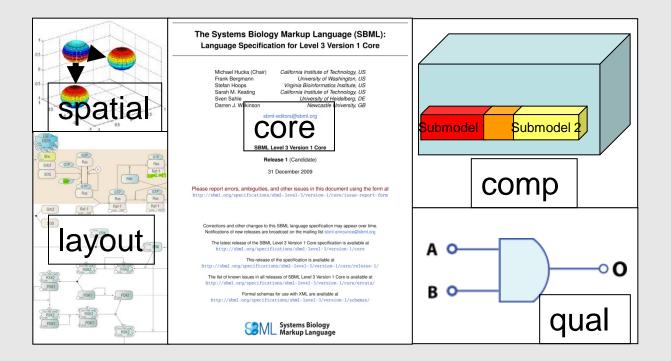






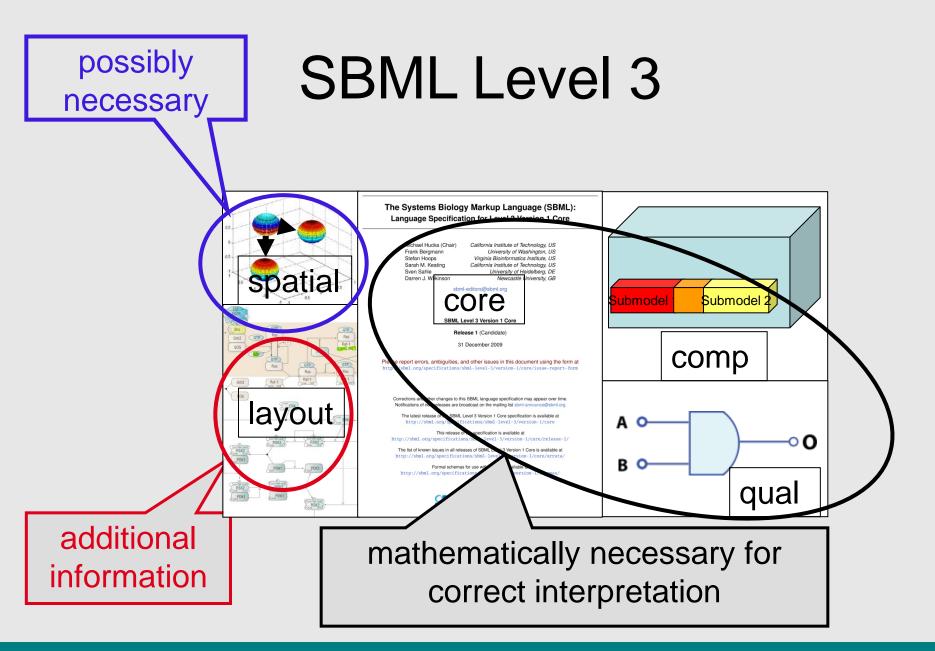
















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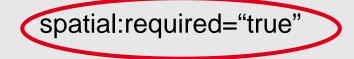
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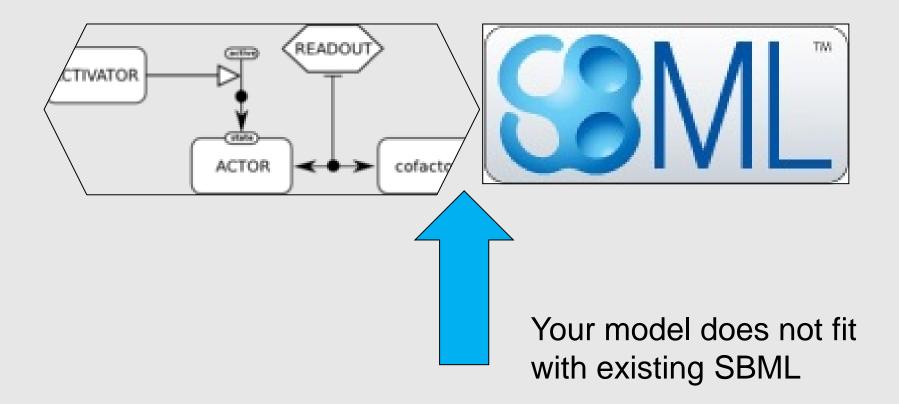
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1. Propose a new package







1. Propose a new package



2. Gather interested parties







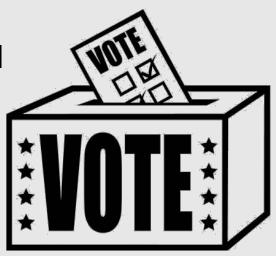




1. Propose a new package

2. Gather interested parties

3. Write a proposal



4. Community vote





1. Propose a new package

2. Gather interested parties

3. Write a proposal

4. Community vote





Voting criteria:

- 1. Address a problem that SBML users would find useful
- 2. Biological processes/phenomena
- 3. Extends SBML naturally from L3 core
- 4. Not a duplication of something existing











Package Working Group

- Mailing list open to anyone
- One or more of the original proposal authors
- One of the current SBML Editors





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Michael Hucka (Chair) Frank Bergmann	California Institute of Technology, US University of Washington, US
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Sven Sahle	University of Heidelberg, DE
Darren J. Wilkinson	Newcastle University, GB
sbm	il-editors@sbml.org
SBML L	Level 3 Version 1 Core
Rel	ease 1 (Candidate)
31	December 2009
	and other issues in this document using the form at sbml-level-3/version-1/core/issue-report-form
	is SBML language specification may appear over time. roadcast on the mailing list sbmi-announce@sbml.org
	Level 3 Version 1 Core specification is available at ications/sbml-level-3/version-1/core
	of the specification is available at ons/sbml-level-3/version-1/core/release-1/
	asses of SBML Level 3 Version 1 Core is available at ions/sbml-level-3/version-1/core/errata/
Formal schema	s for use with XML are available at ations/sbml-level-3/version-1/schemas/

Formal specification



Two separate pieces of software that support the specification







L3 package development Formal specification

- use official SBML Level 3 package template

- include UML diagrams

- provide full explanation





The Systems Biology Markup Language (SBML): Language Specification for Level 3 Version 1 Core Michael Hucka (Chair) Frank Bergmann Stefan Hoops Warbington, US	
Sarah M. Keating California Institute of Technology, US Sven Sahle University of Heidelberg, DE	
Darren J. Wilkinson Newcastle University, GB sbml-editors@sbml.org	
SBML Level 3 Version 1 Core Release 1 (Candidate) 31 December 2009 Please report errors, ambiguities, and other issues in this document using the form at http://sbml.org/specifications/sbml-level-3/version-1/core/issue-report-form	Two so
Corrections and other changes to this SBML language specification may appear over time. Notifications of new releases are broadcast on the mailing list stml-announce@sbml.org	
The latest release of the SBML Level 3 Version 1 Core specification is available at http://sbml.org/specifications/sbml-level-3/version-1/core	
This release of the specification is available at http://sbml.org/specifications/sbml-level-3/version-1/core/release-1/	EXPECT
The list of known issues in all releases of SBML Level 3 Version 1 Core is available at http://sbml.org/specifications/sbml-level-3/version-1/core/errata/	
Formal schemas for use with XML are available at http://shml.org/specifications/shml-level-3/version-1/schemas/	INTERACTION
Systems Biology Markup Language	



Two separate pieces of software that support the specification



Formal specification





Two separate pieces of software that SUPPORT the specification

???

• manipulate constructs

NOT just store and retrieve







Two separate pieces of software that SUPPORT the specification

???

- each tool must manipulate a majority of the constructs
- between them they must manipulate ALL of the constructs

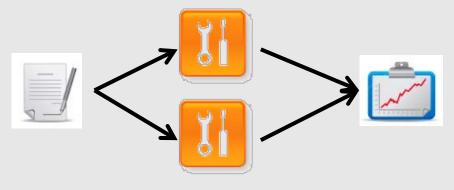




Two separate pieces of software that SUPPORT the specification

???

• test models and corresponding test results

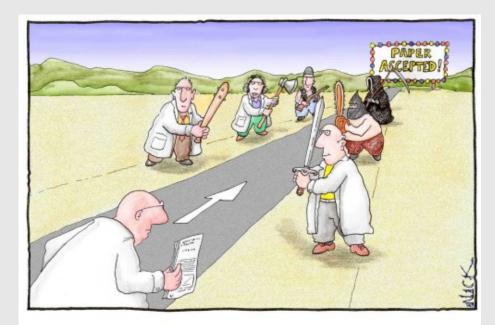




EMBL



SBML Editors review specification and implementations



Most scientists regarded the new streamlined peer-review process as 'quite an improvement.'











L3 package development

Full details of the process:

http://sbml.org/Documents/SBML_Development_Process/ SBML_Development_Process_for_SBML_Level_3





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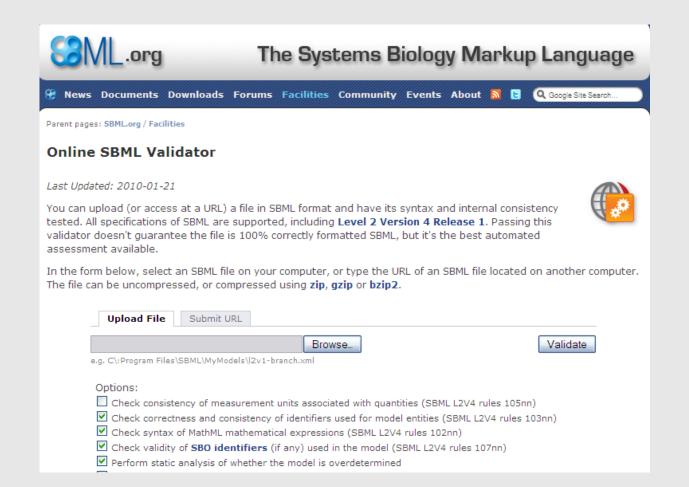
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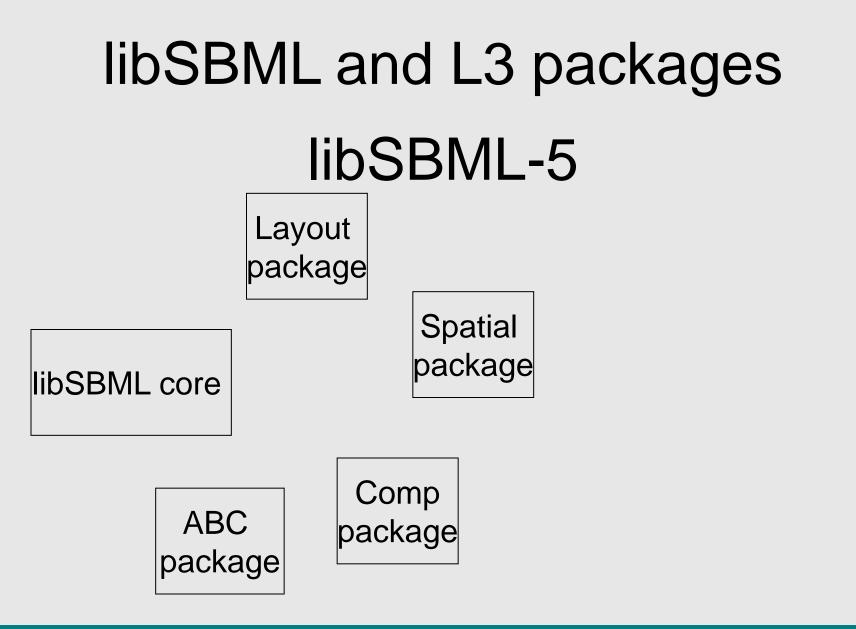
API library for working with SBML

- read
- create
- manipulate
- convert
- write
- validate



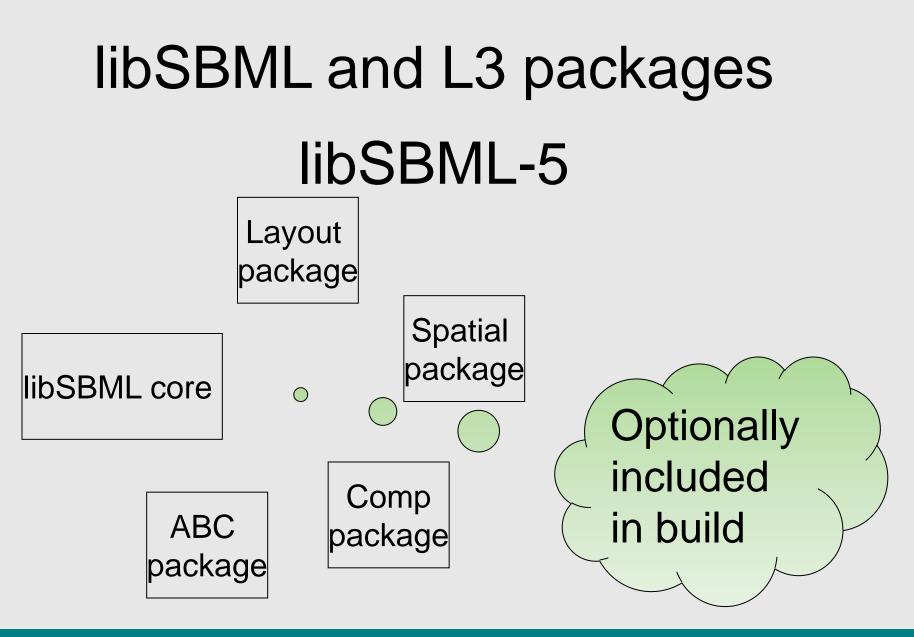
















libSBML and L3 packages

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EXTRA LIBS					





libSBML and L3 packages

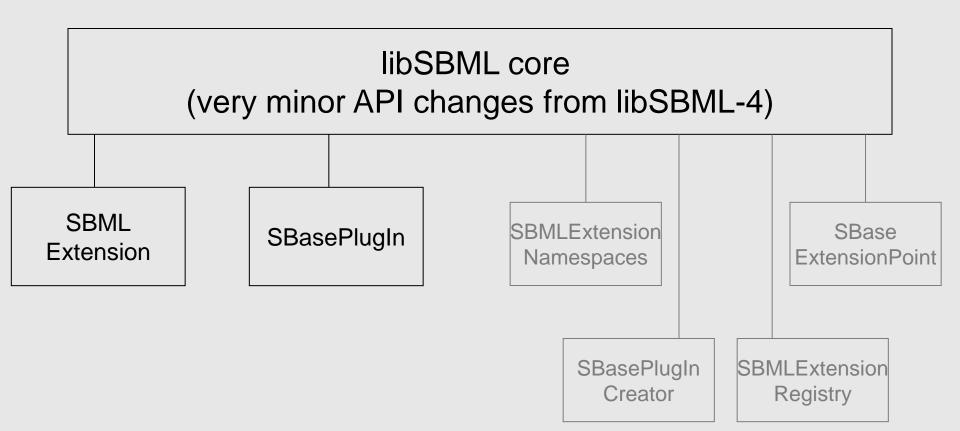
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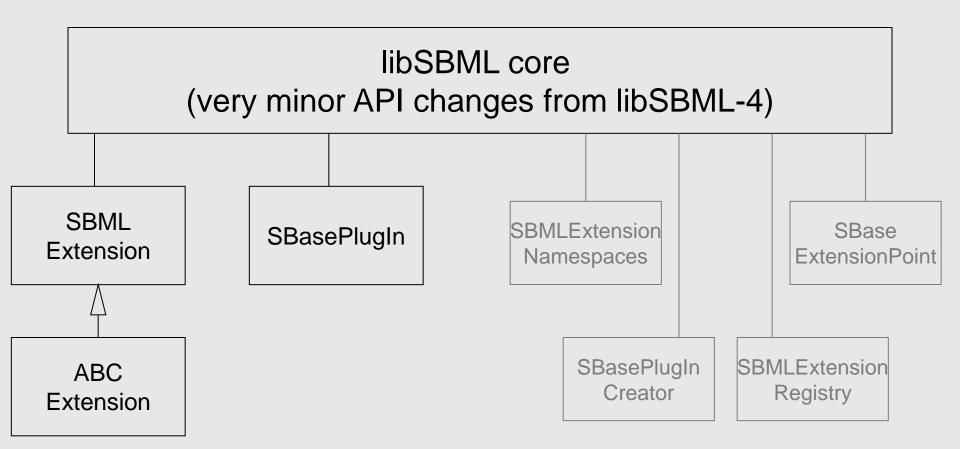
libSBML-5







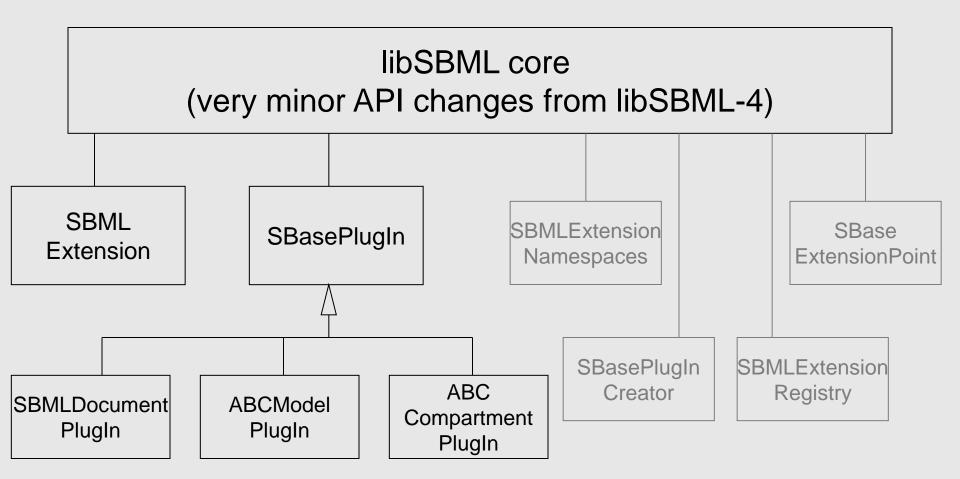
Implementing a package







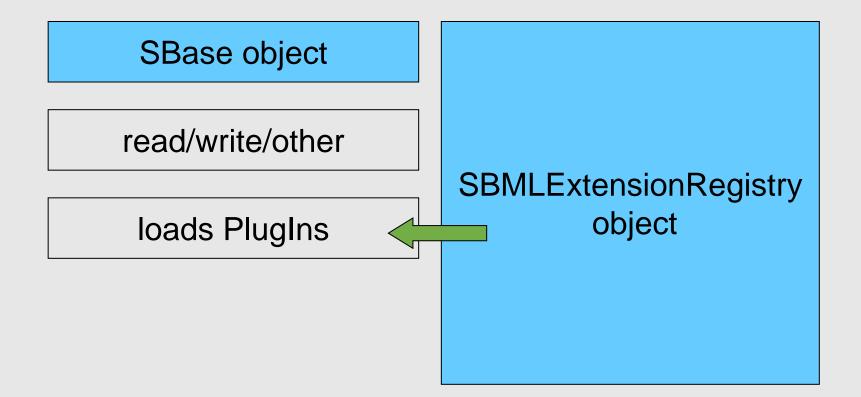
Implementing a package







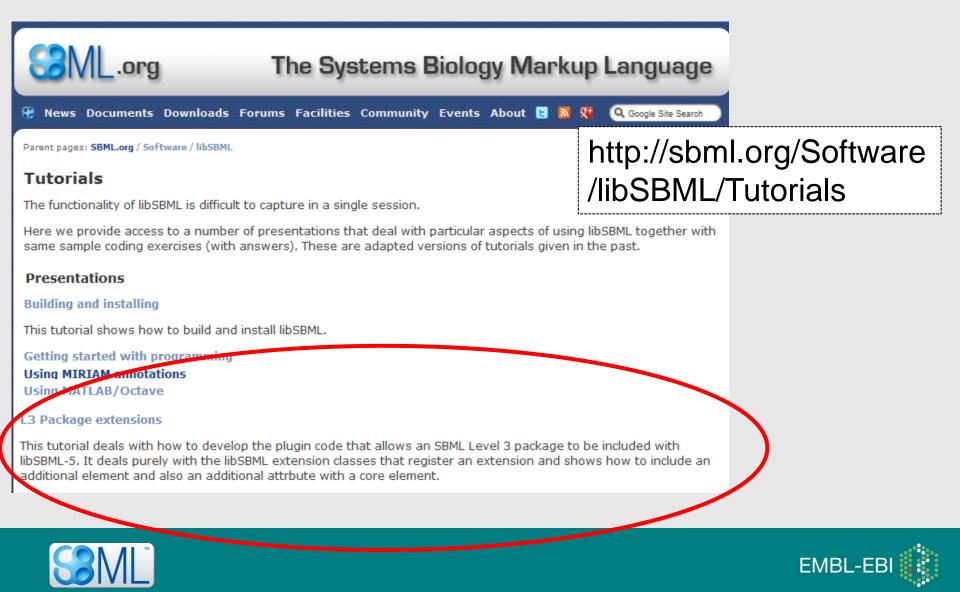
How does it work ?







Implementing a package



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Links

http://sbml.org

Current L3 packages + status: http://sbml.org/Community/Wiki#SBML_Level_3

Process for developing a package: http://sbml.org/Documents/SBML_Development_Process/ SBML_Development_Process_for_SBML_Level_3

libSBML tutorial:

http://sbml.org/Software/libSBML/Tutorials





SBML Team

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John Doyle; Hiroaki Kitano





Acknowledgements

SBML Funding over the years

- NIH National Institute of General Medical Sciences (USA)
- JST ERATO Kitano Symbiotic Systems Project (Japan) (to 2003)
- National Science Foundation (USA)
- International Joint Research Program of NEDO (Japan)
- JST ERATO-SORST Program (Japan)
- Japanese Ministry of Agriculture
- Japanese Ministry of Educ., Culture, Sports, Science and Tech.
- BBSRC e-Science Initiative (UK)
- DARPA IPTO Bio-SPICE Bio-Computation Program (USA)
- Air Force Office of Scientific Research (USA)
- STRI, University of Hertfordshire (UK)
- Beckman Institute, Caltech (USA)
- Molecular Sciences Institute (USA)



