

RepoSE

An On-Line Repository for
Simulation Experiment Descriptions

Maastricht — 21 May 2012



PROBLEMS THAT PLAGUE MODELS

- Same Simulation - Different Results
- Lost Customizations
- Incorrect Initial Values
- Overlooked Procedures
- Broken Functionality



REPRODUCIBLE RESULTS REQUIRE

- Name and version of all the constituent parts
- Detailed description of any changes
- The exact input parameters
- The procedures followed to run the experiment
- List of external dependencies



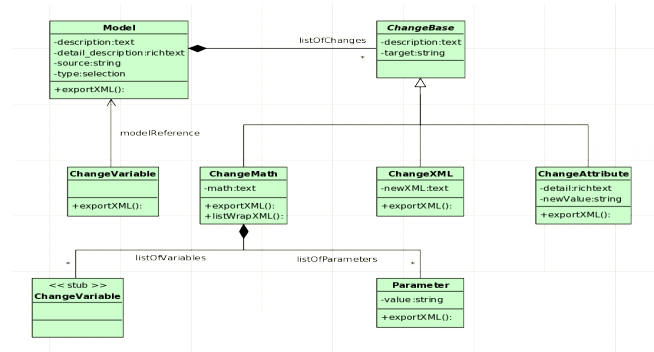
SPECIFY MINIMUM REQUIREMENTS

- Minimum Information Requested In the Annotation of Biochemical Models (MIRIAM)
- Minimum Information About a Simulation Experiment (MIASE)
- Simulation Experiment Description Markup Language (SED-ML)



WHAT IS REPOSE

RepoSE is the realization of the entire SED-ML schema within a content management system named Plone. Each instantiable UML class in the SED-ML schema becomes a content type.



WHAT IS REPOSE'S GOAL

- Make SEDs more understandable by pairing SED-ML's formal precision with prose descriptions
- Make SEDs more accessible by giving modelers control over who can see, append to or edit their SEDs
- Make SEDs more reliable by facilitating curation
- Organize SEDs by the judicial application of generic and domain meta-information



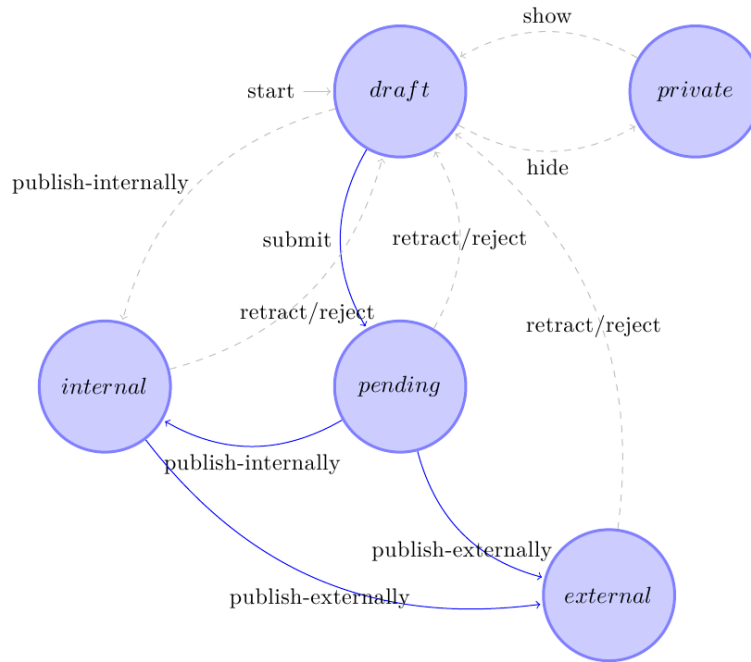
WHAT IS A CONTENT TYPE

- UI components to create, edit and view data
- Machinery that implements the relationships and cardinality of the UML model
- Machinery that reads and writes data from and to the database
- Machinery that complies with Plone's security and workflow systems



WORKFLOWS

Workflows control access and guide curation.



XML BASED SEDs

- RepoSE exports SED-ML compliant simulation experiment descriptions
- SEDs using, e.g., SBML can be exported and run directly



GENERAL PURPOSE SEDs

- We wanted to investigate whether SED-ML could be used as a way to organize simulation experiments written in general-purpose languages. To date we have several SEDs that describe how to build simulation models of developing plant organs using VirtualLeaf.
- Our experience is that models and simulations implemented in code can be of a very different nature where SED-ML concepts do not apply.



THE FUTURE

- Improve the User Interface
- Import existing SED-ML files
- Keep pace with SED-ML development
- Use libSedML to validate and export SED-ML
- Make general purpose SEDs run autonomously



TRY IT YOURSELF

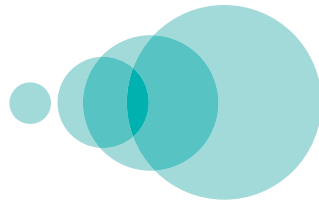
- You'll find the repository at: <http://sed.project.cwi.nl>
- Click on the repository tab
- Choose the SED named O2C2
- Click on the `Export This' link at the bottom right of the page
- Upload and run the SED on the SED-ML Web Tools site.



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NETHERLANDS INSTITUTE FOR
SYSTEMS BIOLOGY



Thank You



Questions?

