

CONNJUR_ML: An XML Schema for NMR Reconstruction Workflows

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Overview

The workflow for modern biomolecular NMR spectroscopy consists of three phases: spectral reconstruction, the process of converting time domain data into the frequency domain; spectral analysis, which includes peak identification and resonance assignment; and biophysical characterization, which includes all subsequent data analysis in which the spectroscopic data is used to draw biophysical inferences (such as structure determination). In this poster we describe an XML schema for describing structural, descriptive and administrative metadata for representing the intermediate datasets generated during spectral reconstruction. As such, this XML schema provides a provenance record of the spectral reconstruction, an essential step in supporting reproducible computation.

PREMIS 3.0 Model

Maintained by the Library of Congress, ***Preservation Metadata: Implementation Strategies*** has been developed in the archives and library communities since 2003 to provide digital preservation systems with a framework to build reliable systems for sustainable information stewardship.

Finalized in 2015, version 3.0 of the PREMIS model extended PREMIS' reach beyond providing administrative and preservation metadata for digital objects to providing for robust, object-oriented, description of physical objects, intellectual entities, and most importantly for this project, computational environmental dependencies.

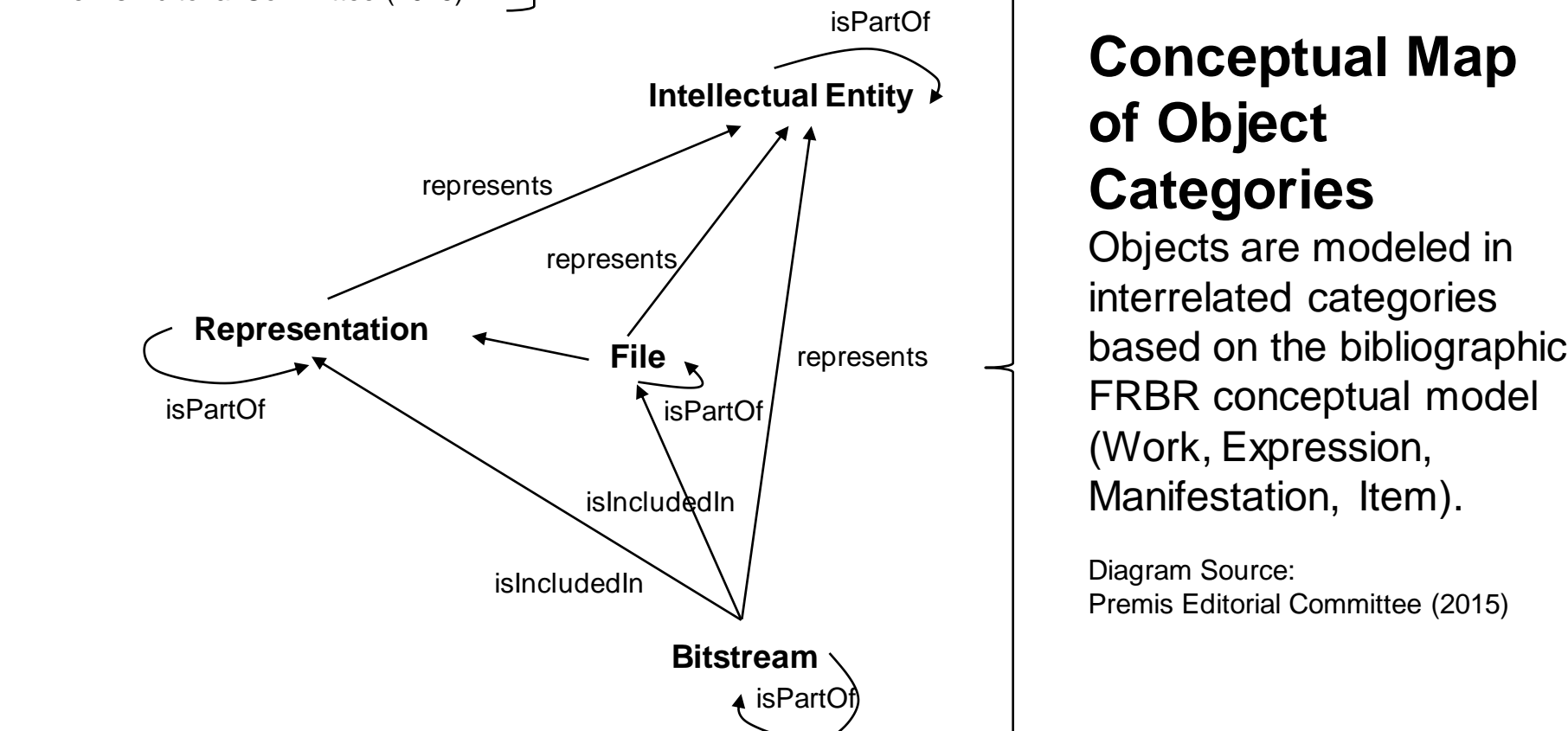
The PREMIS model is encoded as an XML namespace with some RDF semantics.

PREMIS Framework

Semantic Map of PREMIS Objects

Each data-entity is made of several discrete, interrelated objects. CONNJUR_ML focuses on Objects, Events, and Agents. Rights Statements could be used to trigger data access restrictions or data deletion.

Diagram Source:
Premis Editorial Committee (2015)



CONNJUR_ML PREMIS Extension

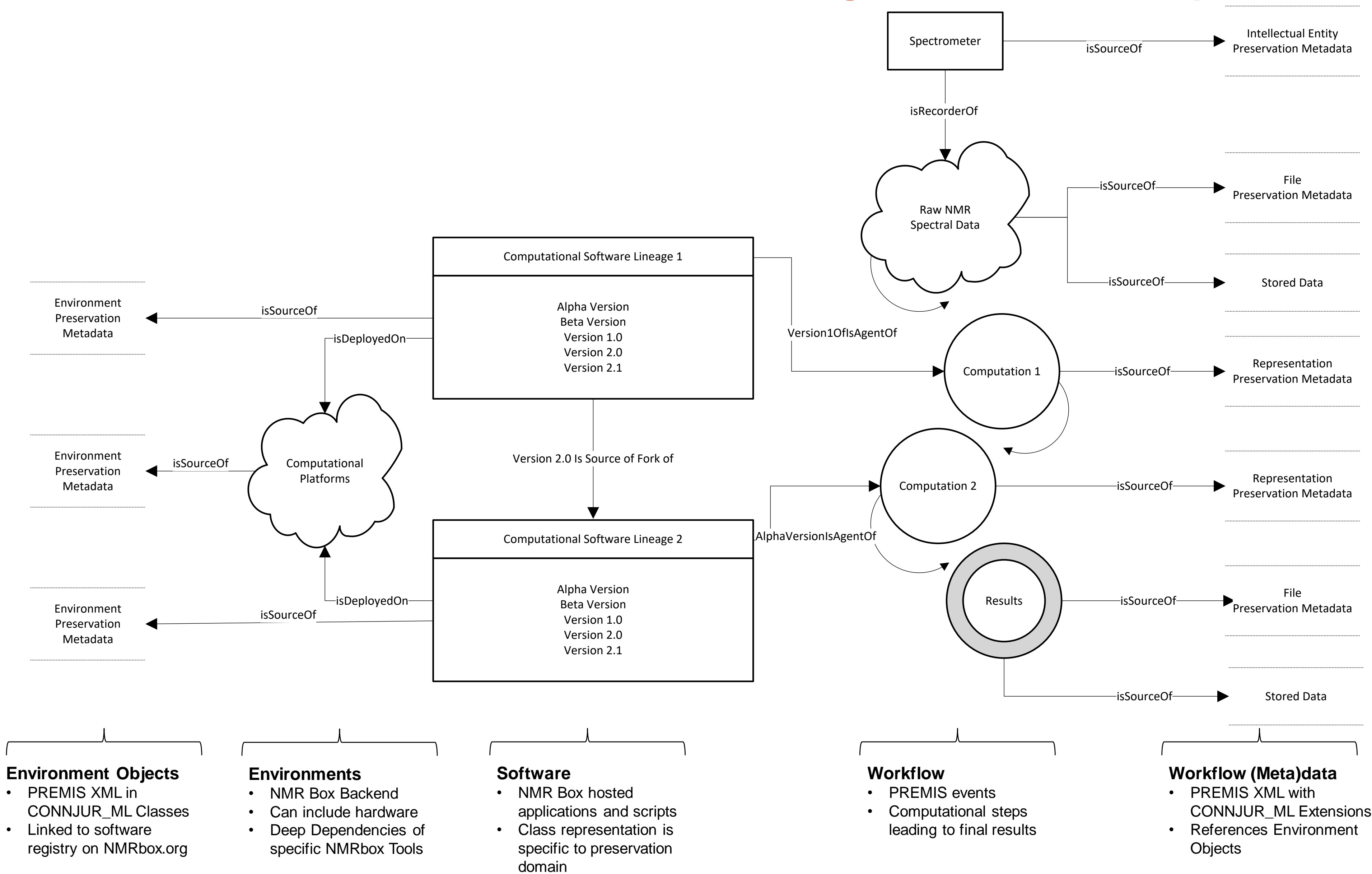
Computational Software Lineages

On ingest into the NMRbox virtualization tool, metadata on software tools are automatically logged as PREMIS objects connect to the NMRbox.org software registry. Related software is encapsulated in classes with group provenance and dependencies that can be accessed from the workflow system.

Self-Referencing Representations as Metadata Surrogates for Ephemeral Pipeline Data

The CONNJUR_ML namespace is embedded in PREMIS *significantPropertiesExtension* when describing workflow metadata for which no intermediate data are stored. Here, metadata transform into the primary data objects that can enhance reproducibility.

Model of a CONNJUR_ML Workflow with Integrated Metadata Capture



Spectral Metadata Embedded in a PREMIS Object with CONNJUR_ML Extensions

```
<?xml version="1.0" encoding="UTF-8"?>
<premis:premis xmlns:premis="http://www.loc.gov/premis/v3"
  xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance"
  xmlns:connjur="https://raw.githubusercontent.com/CONNJUR/CONNJUR_ML/master/connjur_ml.xsd"
  xsi:schemaLocation="http://www.loc.gov/premis/v3 http://www.loc.gov/standards/premis/premis.xsd" version="3.0">
  <premis:object xsi:type="premis:representation">
    <premis:objectIdentifier>
      <premis:objectIdentifierType>local</premis:objectIdentifierType>
      <premis:objectIdentifierValue>NHSQC_001</premis:objectIdentifierValue>
    </premis:objectIdentifier>
    <premis:significantProperties>
      <premis:significantPropertiesType>format</premis:significantPropertiesType>
      <premis:significantPropertiesValue>nmrPipe</premis:significantPropertiesValue>
      <premis:significantPropertiesExtension>
        <connjur:spectrum> <!-- migration -->
          <connjur:spectrumIdentifier>NHSQC_001</connjur:spectrumIdentifier>
          <connjur:spectralFormat>
            <connjur:formatName>nmrPipe</connjur:formatName>
            <connjur:formatSpecification>
              https://www.ihbr.umd.edu/nmrpipe/install.html
            </connjur:formatSpecification>
            <connjur:formatByteOrder>bigEndian</connjur:formatByteOrder>
            <connjur:formatNumericType>IEEE 754-2008 binary32
            </connjur:formatNumericType>
          </connjur:spectralFormat>
          <connjur:spectralDimensions>2</connjur:spectralDimensions>
          <connjur:spectralLayout/>
          <connjur:spectralAxis>
            <connjur:sweepwidth units="Hz">6012</connjur:sweepwidth>
            <connjur:pointType>complex</connjur:pointType>
            <connjur:spacingUnits units="us">166.7</connjur:spacingUnits>
            <connjur:totalPoints>1024</connjur:totalPoints>
            <connjur:sign></connjur:sign>
            <connjur:phaseShift>none</connjur:phaseShift>
            <connjur:encoding>echo-anti-echo</connjur:encoding>
            <connjur:decoupledNucleus>13N</connjur:decoupledNucleus>
            <connjur:constantPhase>0.0</connjur:constantPhase>
            <connjur:linearPhase>0.0</connjur:linearPhase>
            <connjur:domain>time</connjur:domain>
            <connjur:samplingUniform</connjur:sampling>
          </connjur:spectralAxis>
          <connjur:spectralAxis>
            <connjur:sweepwidth units="Hz">2200</connjur:sweepwidth>
            <connjur:pointType>complex</connjur:pointType>
            <connjur:spacingUnits units="us">454.5</connjur:spacingUnits>
            <connjur:totalPoints>256</connjur:totalPoints>
            <connjur:sign></connjur:sign>
            <connjur:phaseShift>none</connjur:phaseShift>
            <connjur:decoupledNucleus>13C</connjur:decoupledNucleus>
            <connjur:constantPhase>0.0</connjur:constantPhase>
            <connjur:linearPhase>0.0</connjur:linearPhase>
            <connjur:domain>time</connjur:domain>
            <connjur:samplingUniform</connjur:sampling>
          </connjur:spectralAxis>
        </connjur:spectrum>
      </premis:significantPropertiesExtension>
    </premis:object>
  </premis:object>
  <!-- ... Additional objects, events, agents and rights ... -->
</premis:premis>
```

Conclusions and Future Work

Implementation Strengths

- Extends 2015 NMR Star Metadata work by Fenwick *et al.*
- Integrates best-practices from information management into scientific workflow
- Creates human readable metadata in a standard archival format with little customization
- XML/RDF allows for mapping and workflow reconstruction with network diagrams

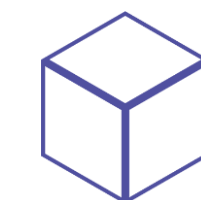
Solicit Feedback

- Understand user needs and potential uses through open dialog with Bio-NMR Community

Future Work

- Middleware automatically generates environment metadata as part of NMRbox registry
- Integration of CONNJUR_ML into existing workflow processes
- Automatic visualization of RDF relationships and network analysis
- Integrate with NMR Star

Acknowledgments



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Repository Access

https://github.com/CONNJUR/CONNJUR_ML



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