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METABOHUB'S SPECTRAL DATABASE: FROM DATA MODEL TO WEB INTERFACE METABOHUB

NILS PAULHE¹, CHRISTOPHE DUPERIER¹, DANIEL JACOB², ÉTIENNE THÉVENOT³, JEAN-FRANCOIS MARTIN⁴, CLAIRE LOPEZ¹, FRANCK GIACOMONI¹

BOLISM EXPLORATION PLATFORM, INRA (CLERMONT-FERRAND/THEIX) ² BORDEAUX METABOLOME PLATFORM, INRA (BORDEAUX) ³THE METABOLOMEIDF, CEA (SACLAY) ⁴ METABOLOMICS AND FLUXOMICS PLATFORM, INRA (TOULOUSE)

Introduction

MetaboHUB is an infrastructure of metabolomics and fluxomics that provides tools to research teams and partners; a full workpackage focuses on bioinfomatics tools design and development, including one called "MetaboHUB's Spectral Database". This tool is created to gather and share knowledge and data from both Mass and NMR Spectra.

The core of the database called "data-model", is a computational representation of each entity involved in spectra analysis and chemical compounds annotation.

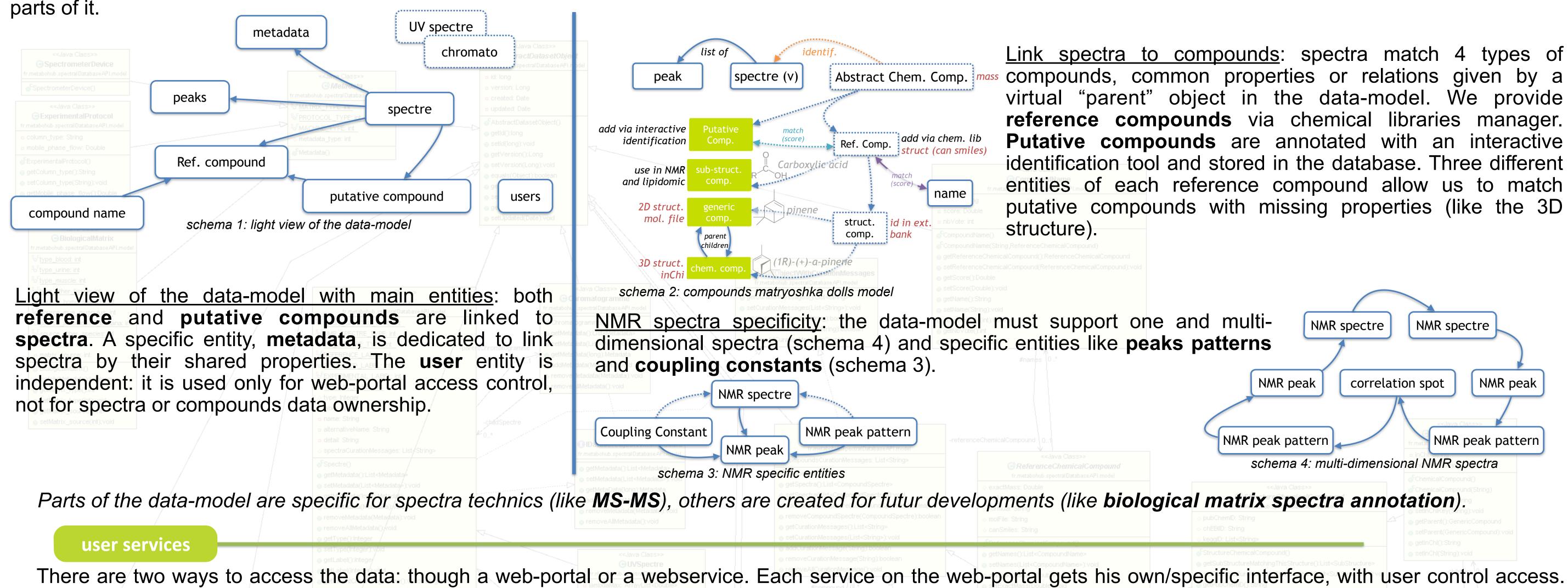
It was designed with spectrometry experts knowledge; their experience will also be required to build an interactive identification interface for spectral analysis. This database will be available through a web-portal and open for each partner of the MetaboHUB consortium in a first time, to the whole community later.

Methods

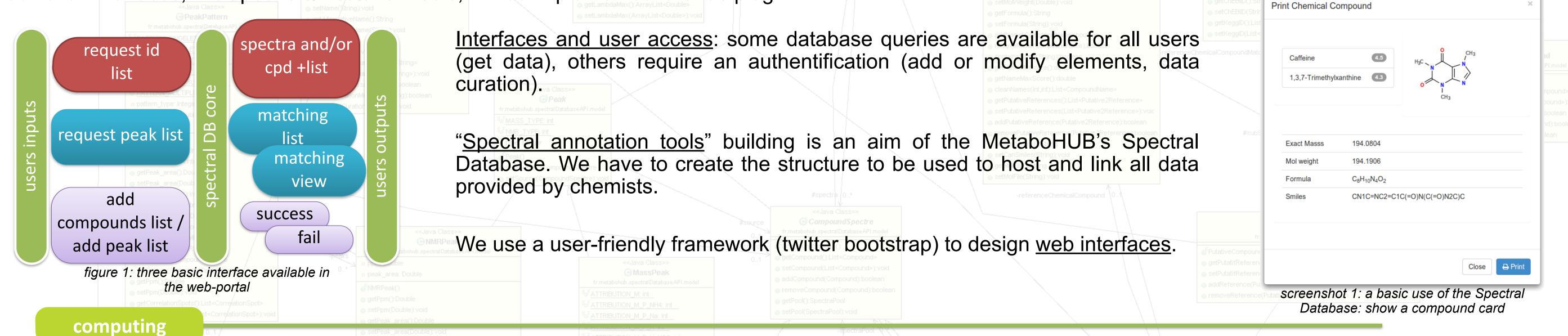
For the MetaboHUB's Spectral Database code architecture, we use modelling concepts and objects-objects relations to represent mass / nuclear magnetic resonance (NMR) spectrometry complexity. Manipulated entities and their links are based on data / metadata used by chemists in laboratory routine. Our work is split in three levels with data-model building, users services designing and code engineering.

data model

Removing the complexity of the full data-model allows us a better understanding of it; the first schema proposes a light view of relation entities, others are focused on







Code architecture and frameworks are an originality of this project. Developers use java frameworks and common generic toolbox to keep to the time scheduled and interoperability. There are three levels of architectures: the spectral database core (figure 2), implemented in both webservices and the web-portal model-view-controller (MVC) (schema 5). load/store database

<u>Code architecture</u>: the MVC pattern is better for code maintainability, team development and security.

<u>Webservices</u> are the way to open with bots data stored in the spectral database, submitting massives requests and hudge computing. We use popular technologies like REST and Json.

core	ext. resources connectors external-tools io-galaxy external-banks
	files parser io-spectra-files io-chem-files data model
	figure 2 : core of the Spectral Database

The core of the database is split in stand-alone blocks, which allow the community to re-use them in others projects. Each block is specific to one function.

To store data in the database, we use the Hibernate framework. We keep the model relations and constrains, with an abstraction of technologies used for SQL requests.



Conclusion

Still in early development phase, some functional blocks are already being tested.

The first challenge was to validate this data-model and the use cases with chemists. We are currently testing the chemical library module with data provided by MetaboHUB partners.

Two milestones are coming: a first to provide a mechanism to import spectral data in the data-model and feed the database, a second to define metadata around spectral analysis.



core

manipulates

controller

uses 📃

security

mode

user

updates

view

schema 5: MVC structure

user

friendly



INRA - Unite de Nutrition Humaine Plate-Forme "Exploration du Métabolisme" Centre de Theix 63122 Saint genes Champanelle France



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