Metabolomics Resources in the Biological Magnetic Resonance Data Bank (BMRB)

Kumaran Baskaran, Jonathan R. Wedell, Hongyang Yao, Dimitri Maziuk, Hamid R. Eghbalnia, Michael M. Gryk, and Jeffrey C. Hoch

Department of Molecular Biology and Biophysics, UConn Health, 263 Farmington Avenue, Farmington, CT 06030, USA
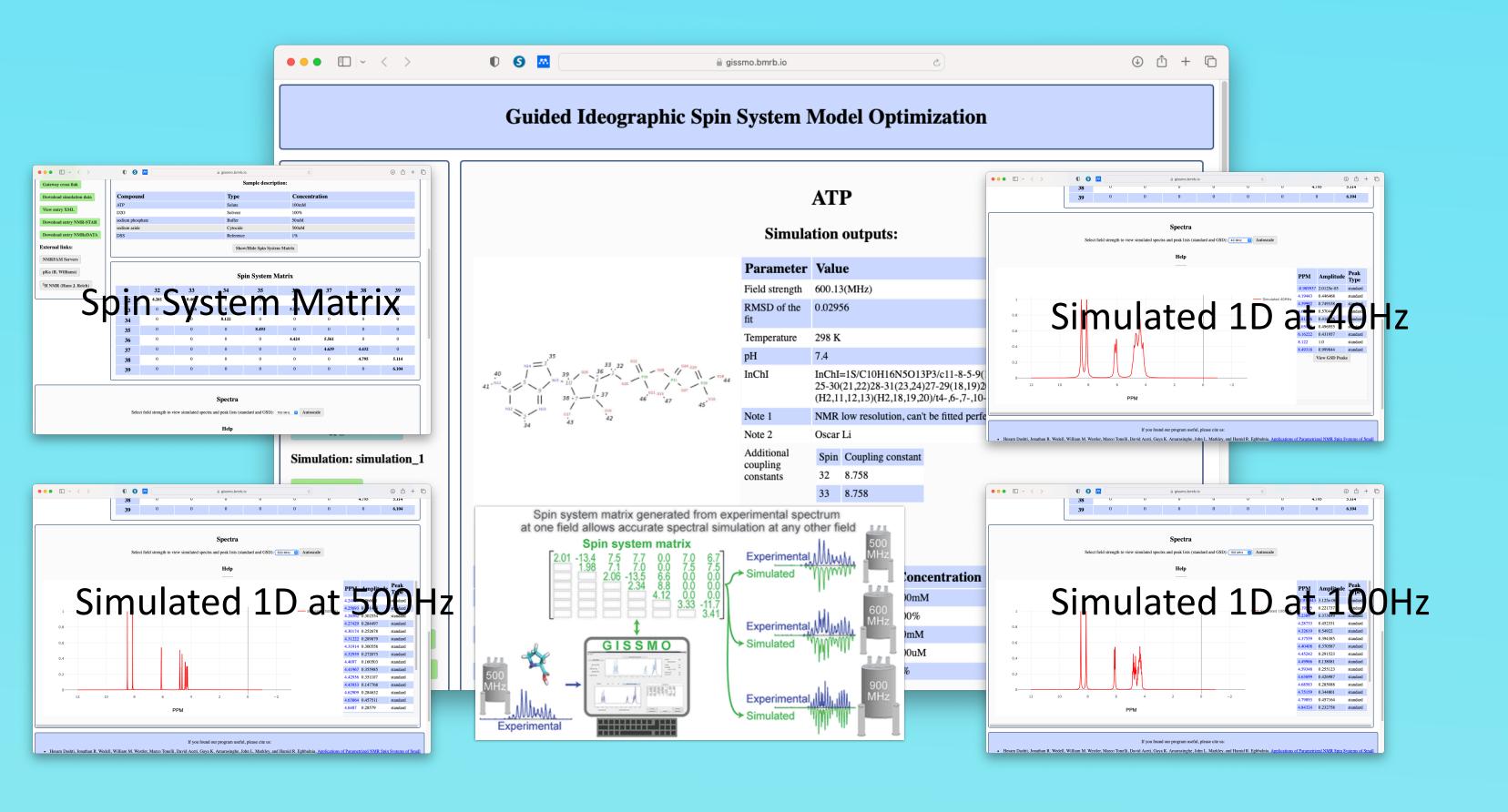
BMRB small molecules library
(https://bmrb.io/metabolomics/)

BMRB[1] provides curated set of reference NMR spectroscopic data for more than 1000 metabolites measured using a standardized experimental protocol

Each reference data set consists of
- Chemical shift assignments
- Sufficient time domain data (FID) from a set of 1D and 2D experiments to analyze the molecule

This library could be used to analyze and identify molecules in bio-fluids and other metabolite mixtures. Peak lists of 1D and 2D experiments can be searched across the database through web interface or using BMRB-API
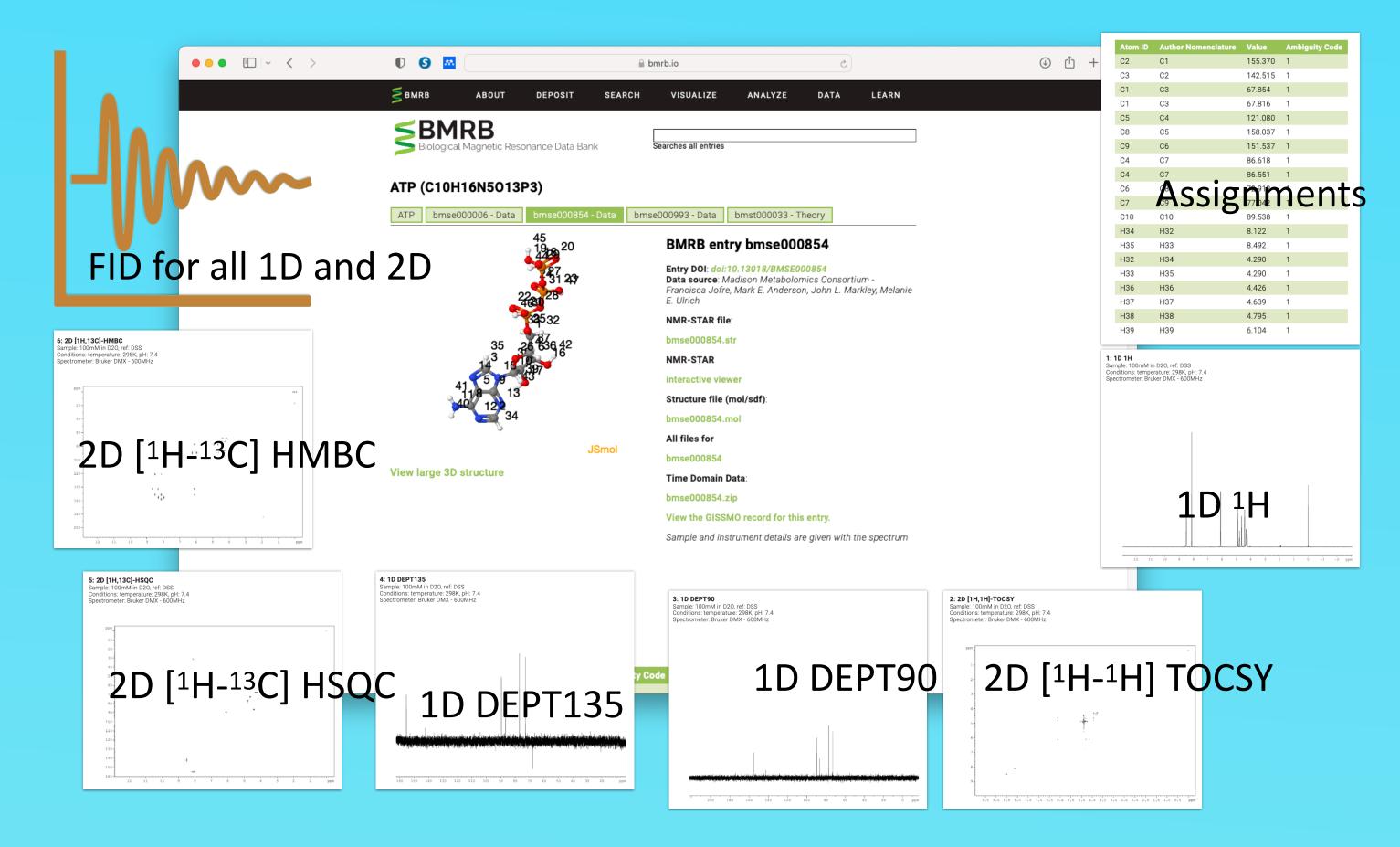

GISSMO Library
(https://gissmo.bmrb.io/)

Guided Ideographic Spin System Model Optimization[2](GISSMO) enables the efficient calculation and refinement of spin system matrices(chemical shift and coupling constants against experimental 1D-^{1}H spectra of small molecules.

GISSMO library includes
- Library of over 1000 parameterized small molecules
- Spin system matrix for every molecule
- Simulated 1D spectra at various field strength (40Hz to 1.3GHz) and peak lists at each field strength

ALATIS Library
(https://alatis.bmrb.io/)

Atom Label Assignment Tool Using InChI String[3] (ALATIS) creates unique InChI identifiers for small molecules through rigorous labeling of their atoms.This numbering system helps to identify molecules across different databases and flawlessly compare the properties of their individual atoms

ALATIS library includes
- Unique InChI strings for molecules from various databases like BMRB, PubChem, HMDB and RCSB-PDB Ligand-Exp
- Cross-links from PDB entries to BMRB, HMDB and PubChem

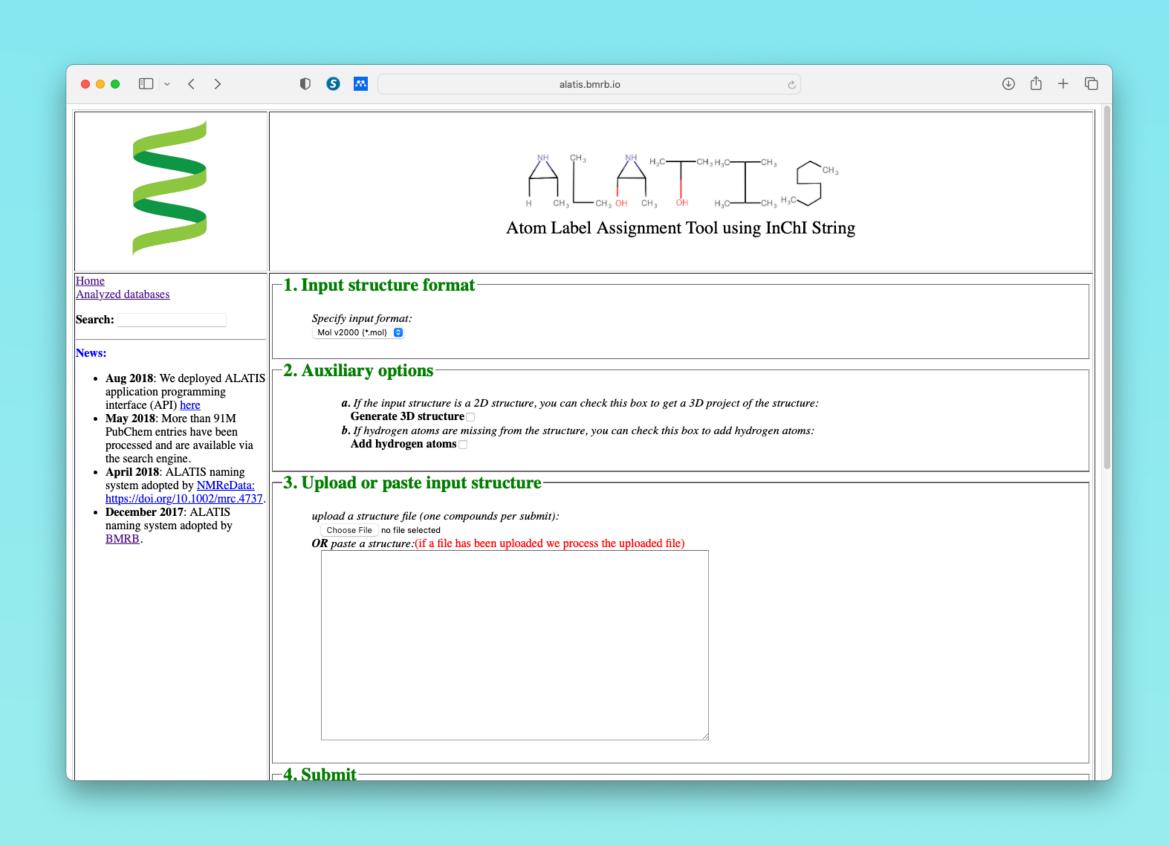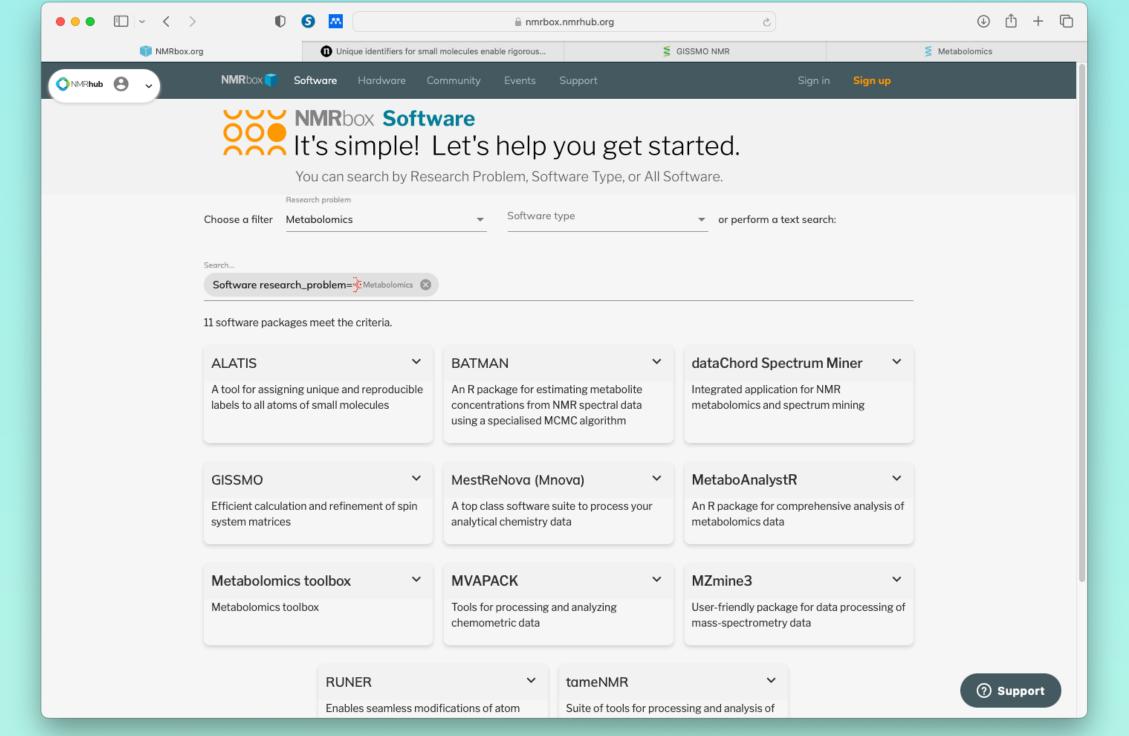BMRB follows ALATIS numbering system in its small molecules database.

NMRbox: Software and Computing resource
(https://nmrbox.nmrhub.org/)

NMRbox[4] is a resource for biomolecular NMR software. It provides tools for finding the software you need, documentation and tutorials for getting the most out of the software, and cloud-based virtual machines for executing the software. NMRbox is available free for academic use.

NMRbox provides ready to use computing resource along with various preinstalled software tools like **ALATIS, BATMAN, dataChord Spectrum Miner, GISSMO, MestReNova(Nova). MetaboAnalystR, Metabolomcis toolbox, MVPACK, MZmine3, RUNNER, tameNMR** and **MATLAB**.
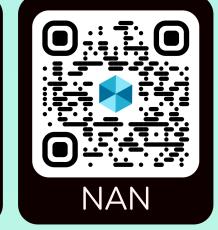
NMRHhub is an umbrella organization which grants access to data collection infrastructure (Network of Advanced NMR: NAN), analysis platform (NMRbox), data resource (BMRB) and Non-uniform sampling and reconstruction project (NUScon) through single portal.
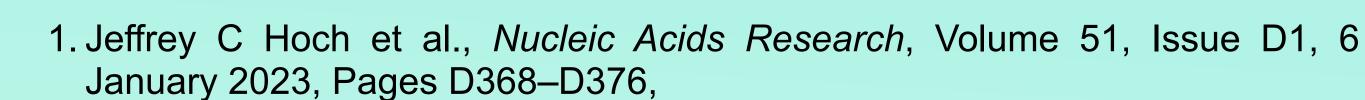

NMRbox   NUScon   NAN   BMRB

1. Jeffrey C Hoch et al., *Nucleic Acids Research*, Volume 51, Issue D1, 6 January 2023, Pages D368–D376,
2. Hesam Dashti et al., Analytical Chemistry 2018 90 (18), 10646-10649
3. Hesam Dashti et al., Sci Data 4, 170073 (2017)
4. Jeffrey C Hoch et al.., Biophysical Journal, 112: 1529-1534, 2017.