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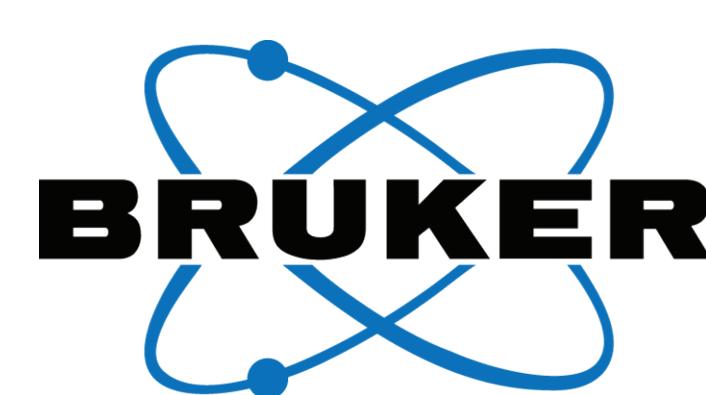
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Magnetic Resonance in Chemistry, Wiley, Chichester, UK

Editor in Chief

Member of the Associate board

Executive journal editor



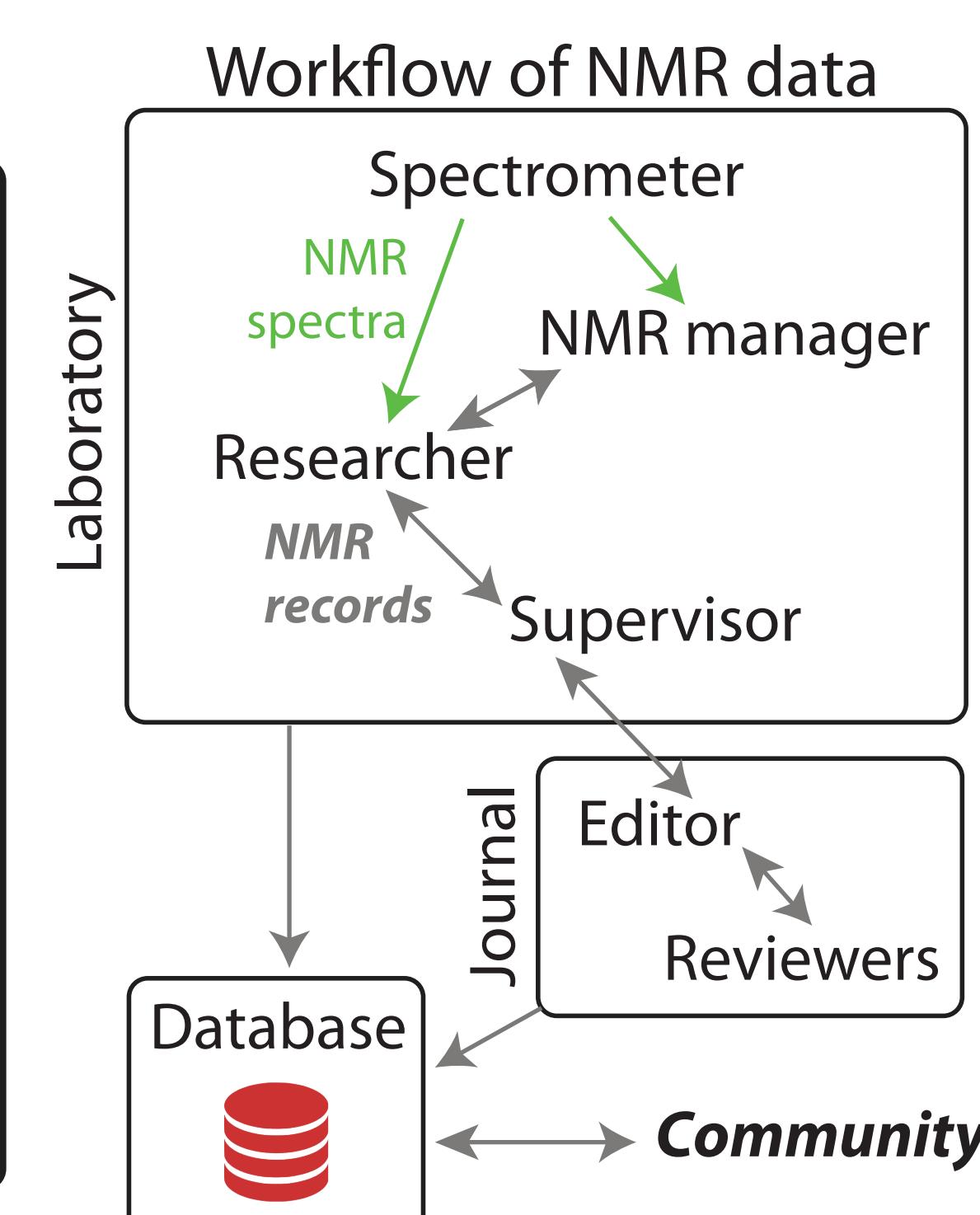
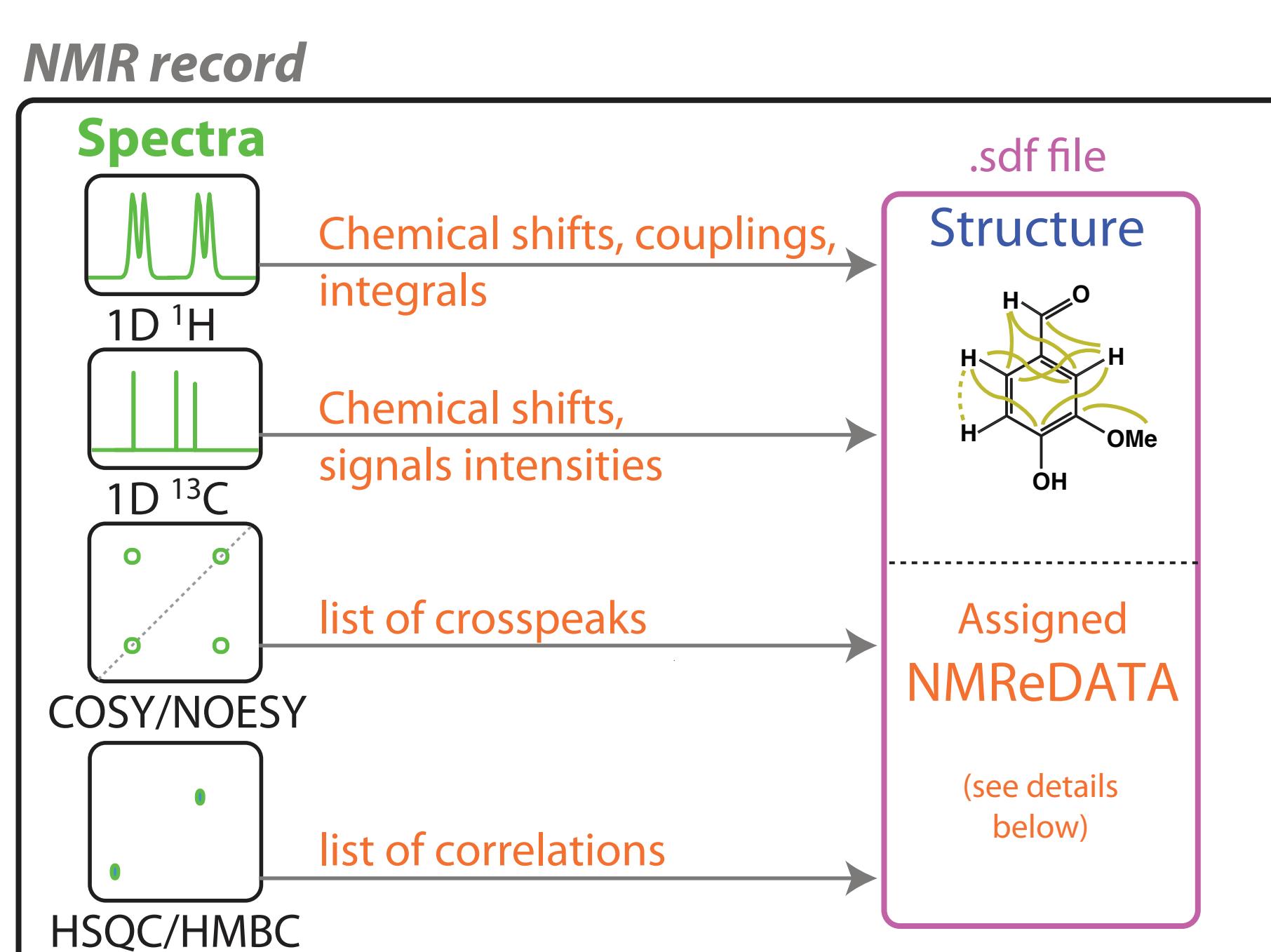
The NMReDATA initiative

The goal of the NMReDATA initiative is to introduce a manner to associate the data extracted from a "full" NMR analysis (the NMReDATA) to a chemical structure.

We introduced a file format based on the commonly used "Structure Data Format" (.sdf) to combine the chemical shifts, couplings, lists of 2D correlations and assignment (NMReDATA) with a chemical structure in the .mol format.

NMR records (NMR spectra + NMReDATA) including the .sdf file will be generated by computer-assisted structure elucidation software or web-based tools under development.

D. Jeannerat, Magn. Res. in Chem., 2017, 55, 7-14.
J. Bisson, C. Simmler, S.-N. Chen, J. B. Friesen, D. C. Larkin, J. B. McAlpine, G. F. Pauli, Nat. Prod. Rep., 2016, 33, 1028



Important benefits of the new format

- Improved quality of the NMR data for researchers and the community
- Straightforward inclusion of NMR data in reports and journal articles
- Simplified referee work
- Compatibility with electronic storage in database
- Easier comparison of dataset
- Improved searchability of NMR data

Detailed structure of NMR records

```
benzo[a]pyrene.nmredata.zip
benzo[a]pyrene
  10 (spectrum 1)
  11 (spectrum 2)
  12 (spectrum 3)
  13 (spectrum 4)
benzo[a]pyrene.sdf

"structure" part of the file
The format is the same as .mol files (compatible with most chemical structure editor)
list of atoms with coordinates (either 2D as in "flat" structures used for drawing or 3D)
list of bonds
Explicit structures (including all hydrogen atoms) should be favored to facilitate the assignment of non-equivalent hydrogens. For aromatic compounds implicit hydrogen is fine.

"NMReDATA" part of the file
The .sdf file can include "tags". Each tag has a name between ">" <" and >" and data (one or more lines terminating with an empty line).
NMReDATA tags start with "NMREDATA_".
  0.96
  <NMREDATA_ID>
  no id yet
  <NMREDATA_SOLVENT>
  cdc13
  <NMREDATA_TEMPERATURE>
  298.15
  <NMREDATA_SIGNALS>
    This tag associates the labels used in the assignment and the atom(s) of the molecule.
    Labels used for the assignment (see right column). In this example, the proton bound to the atom 1 in the structure part (in blue above) was called "H(1)" and the carbon "(1)", but the authors can chose the labels according to IUPAC rules or the requirement of the journal where the data are published.
    Chemical shifts in ppm
    Atom number in the .mol part (see the blue frame above). When using implicit hydrogen, the atom number to which the hydrogen atom is bound is given following "H".
  H(1), 128.17, 1
  (2), 127.9, 2
  (3), 129.4, 3
  (4), 131.6, 4
  (5), 123.8, 5
  (6), 124.4, 6
  (7), 129.9, 7
  (8), 128.3, 8
  (9), 127.4, 9
  (10), 131.35, 10
  (11), 126.09, 11
  (12), 131.4, 12
  (13), 128.23, 13
  (14), 122.2, 14
  (15), 125.6, 15
  (16), 127.5, 16
  (17), 125.9, 17
  (18), 126.9, 18
  (19), 125.97, 19
  (20), 126.06, 20
  H(1), 8.02, H1
  H(2), 7.94, H2
  H(6), 8.53, H6
  H(7), 8.10, H7
  H(11), 7.98, H11
  H(14), 9.07, H14
  H(15), 8.25, H15
  H(16), 7.95, H16
  H(19), 7.85, H19
  H(20), 7.79, H20
  H(17), 9.06, H17
  H(18), 8.30, H18

  > <NMREDATA_1D_1H>
  ;note that integrals (E) were not measured in the spectrum but set to 1
  Spectrum_Location=file://nmr/10/pdata/1/
  Larmor=500.13
  Decoupled=1H
  100, I=100.00
  131, I=100.00
  131, I=100.00
  129.9, I=100.00
  128.9, I=100.00
  128.23, I=100.00
  127.9, I=100.00
  127.4, I=100.00
  127.5, I=100.00
  126.09, I=100.00
  126.09, I=100.00
  126.09, I=100.00
  125.6, I=100.00
  125.6, I=100.00
  124.8, I=100.00
  124.9, I=100.00
  123.8, I=100.00
  123.0, I=100.00
  122.2, I=100.00
  Spectrum_Location=file://10/1/pdata/1/
  > <NMREDATA_1D_13C>
  ;note that intensities were not measured in the spectrum 110 for CH 100 for Cg
  Spectrum_Location=file://nmr/13/pdata/1/
  Larmor=125.0
  Decoupled=1H
  100, I=100.00
  131, I=100.00
  131, I=100.00
  129.9, I=100.00
  128.9, I=100.00
  128.23, I=100.00
  127.9, I=100.00
  127.4, I=100.00
  127.5, I=100.00
  126.09, I=100.00
  126.09, I=100.00
  126.09, I=100.00
  125.6, I=100.00
  125.6, I=100.00
  124.8, I=100.00
  124.9, I=100.00
  123.8, I=100.00
  123.0, I=100.00
  122.2, I=100.00
  Spectrum_Location=file://13/1/pdata/1/
  > <NMREDATA_2D>
  Isotope in F1 / type of mixing / isotope in F2
  > <NMREDATA_2D_1H_NJ_1H>
  Larmor=500.13
  CorType=COSY
  H(1)/H(2)
  H(7)/H(11)
  H(11)/H(15)
  H(14)/H(16)
  H(19)/H(20)
  H(19)/H(17)
  H(20)/H(18)
  Spectrum_Location=file://12/1/pdata/1/
  > <NMREDATA_2D_13C_1J_1H>
  one-bound correlations test of command line
  Larmor=500.13
  CorType=HSQC
  (1)/H(1)
  (2)/H(2)
  (6)/H(6)
  (7)/H(7)
  (11)/H(11)
  (14)/H(14)
  (15)/H(15)
  (16)/H(16)
  (17)/H(17)
  (18)/H(18)
  (19)/H(19)
  (20)/H(20)
  Spectrum_Location=file://13/pdata/1/
  ... HMBC, NOESY data would follow with the same format
  > <NMREDATA_LITERATURE>
  Source=Journal
  DOI=...
  > <...>
  Reference to the main publication associated to the record.
  Other data such as the origin of the sample, the sample preparation and other analytics data could be added to .sdf file outside the NMReDATA initiative.
  $$$$ End of file code
```

Timeline of the initiative

Mid-2016: Proposition by the members of the Associate editorial board of Magnetic resonance in Chemistry to request authors to submit NMR spectra and the extracted data in a manner allowing serious reviewing and to become a source or reliable peer-reviewed NMR data.

September 2016: Decision of the Editorial board of Magnetic Resonance in Chemistry to request NMR data for structure papers.

Until March 2017: Elaboration of a beta version of the format to include NMR data in .sdf files.

July 2017: Official announcement of the Initiative at the Euromar 2017, Warsaw, Poland.

By August 2017: Elaboration and tests of the cheminfo.org web-based platform to host the embargoed NMR records during the peer-review process and open them to the public upon acceptance for publication.

By September 2017: Implementation of import/export features by the providers of computer-assisted structure elucidation software.

September 2017: Round-table discussion at the SMASH conference (Baveno, Italy) and decision on the version 1.0 of the file format.

From January 2018: NMReDATA and spectra will be requested for all structure papers submitted to Magnetic Resonance in Chemistry.

Mid-2018: Evaluation of the initiative. If satisfactory, invitation to other journals to join the initiative.

During 2018: Contact with IUPAC to consider changes in the recommendation to report small-molecule NMR data.

Follow the progress of the initiative on
www.nmredata.org



Acknowledgments

Damien Jeannerat thanks the State of Geneva and the Swiss NSF funds 200021-147069, 200020-126650, 206021-1288746 for financial support. This work does not necessarily reflect U.S. EPA policy.