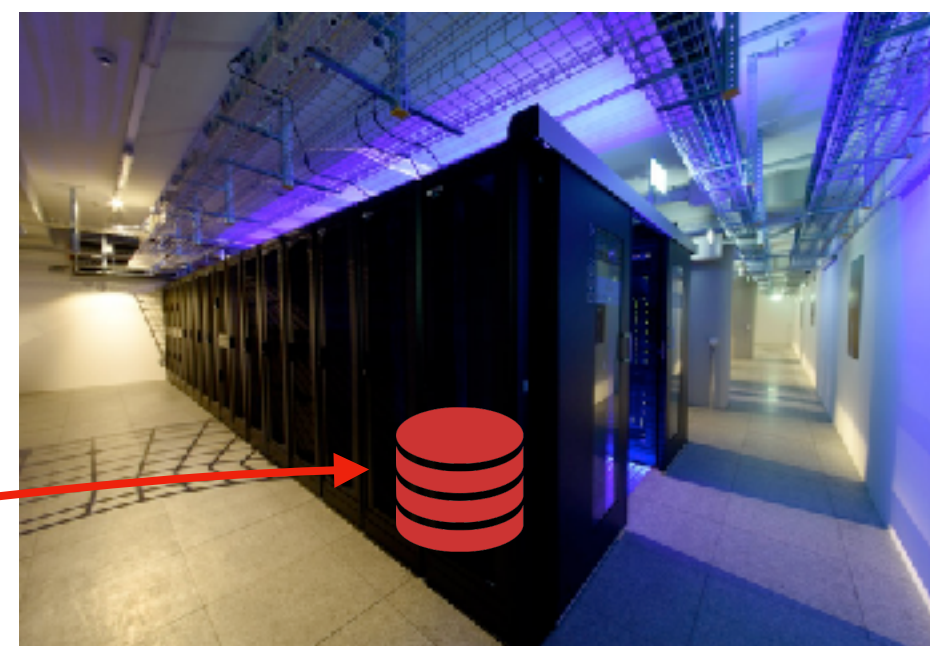
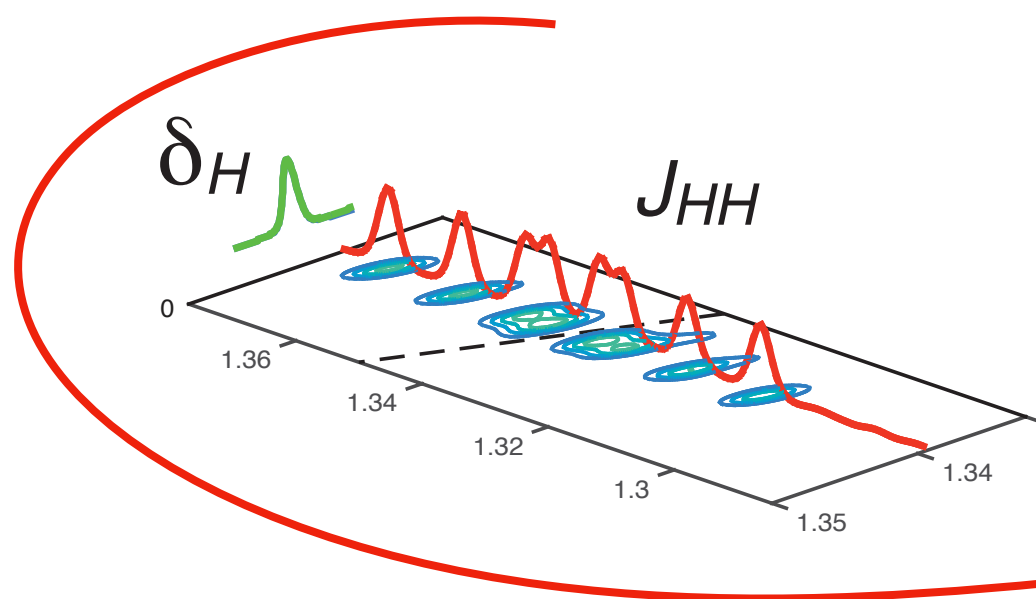
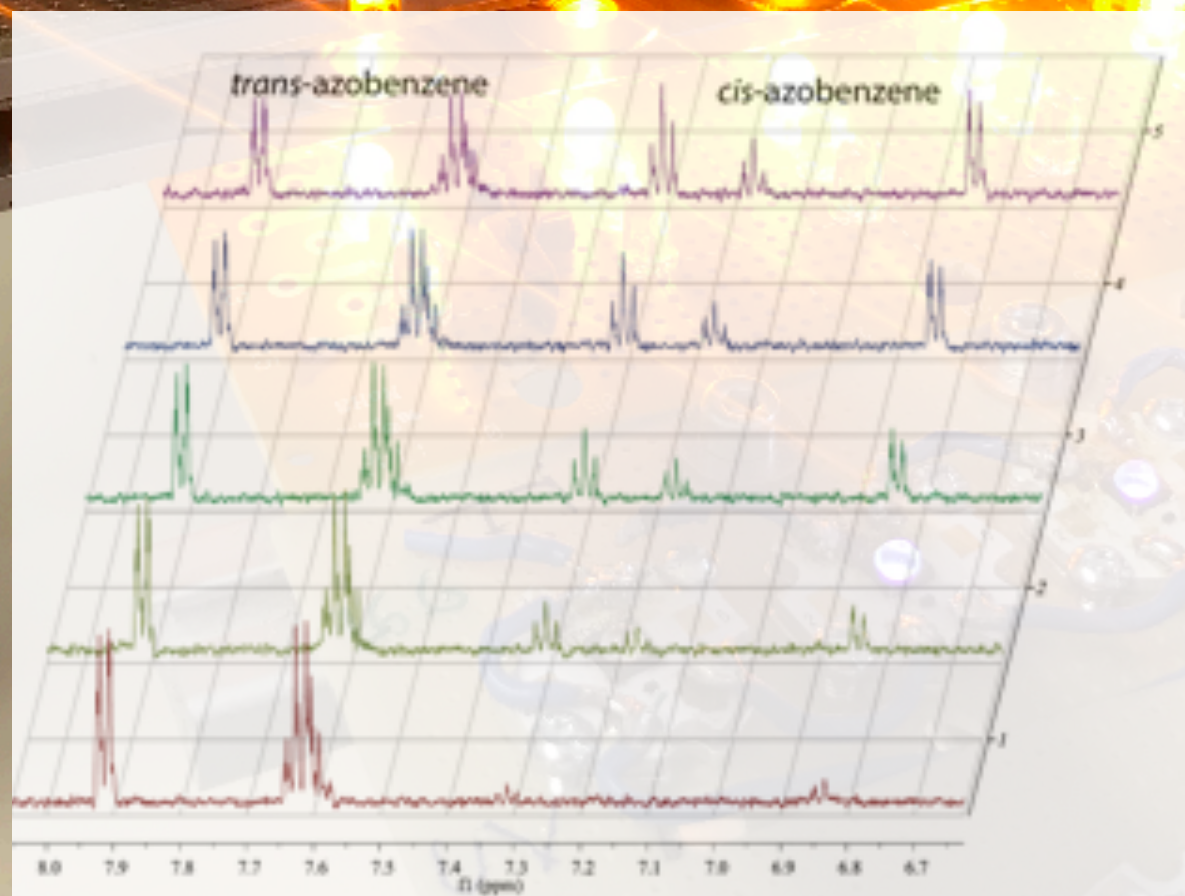
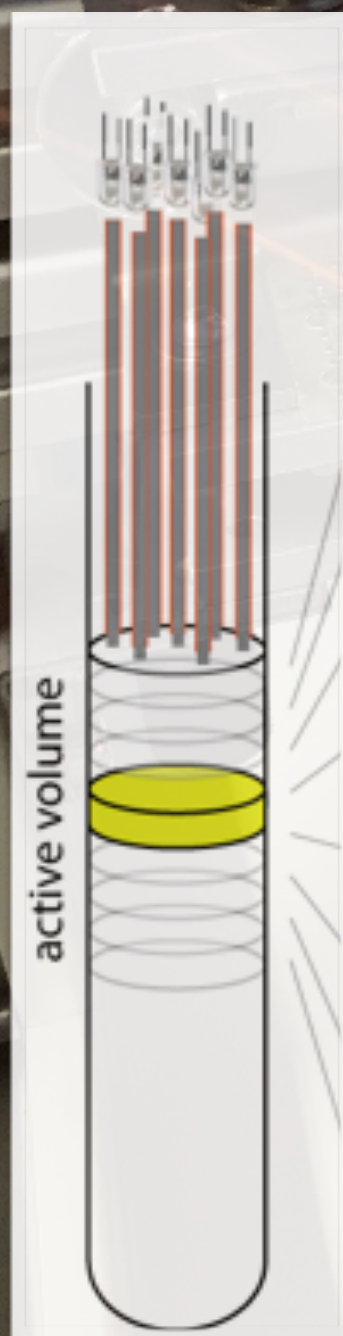
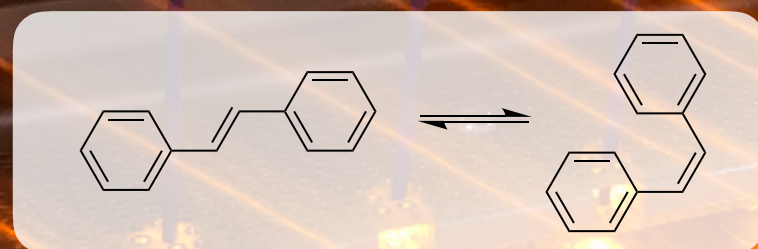
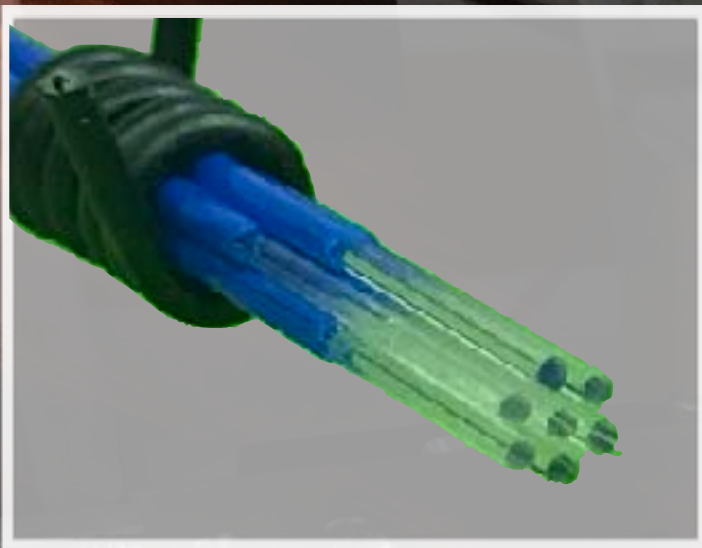


NMReDATA: a format and a good practice to exchange NMR parameters associated to chemical structures



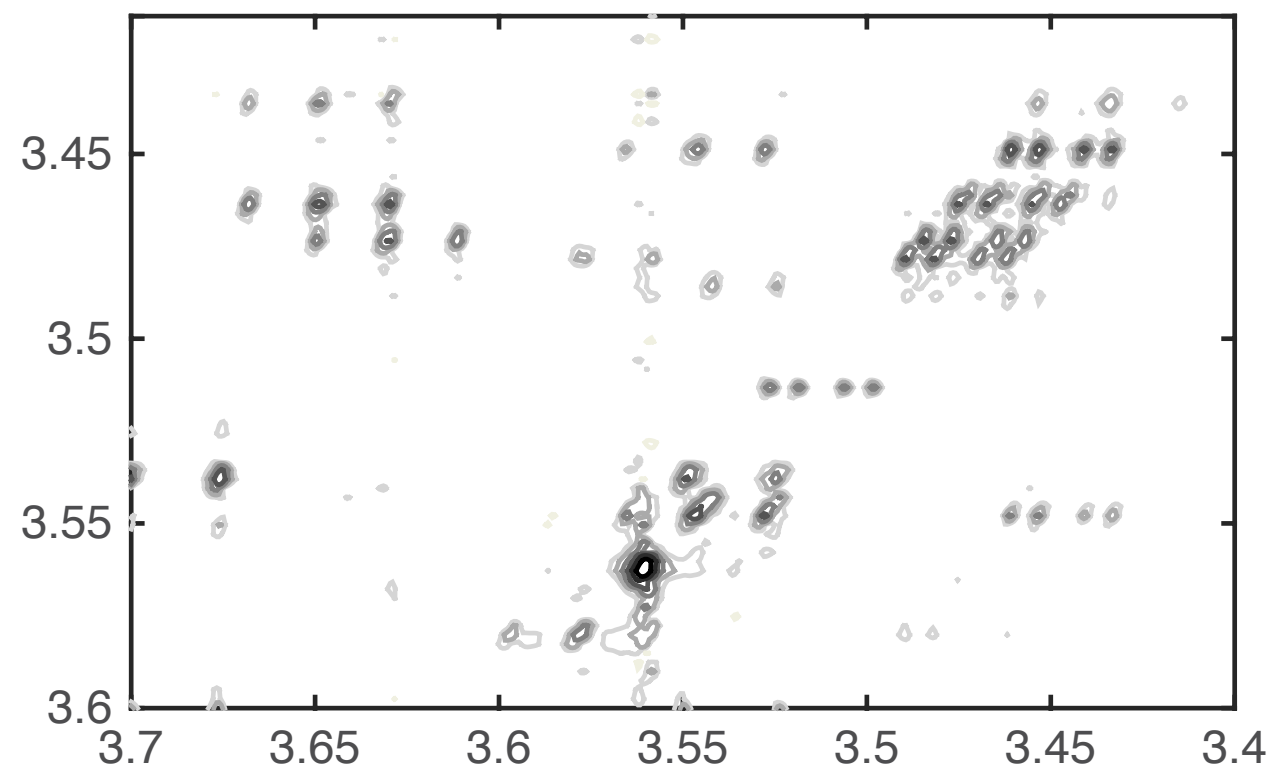
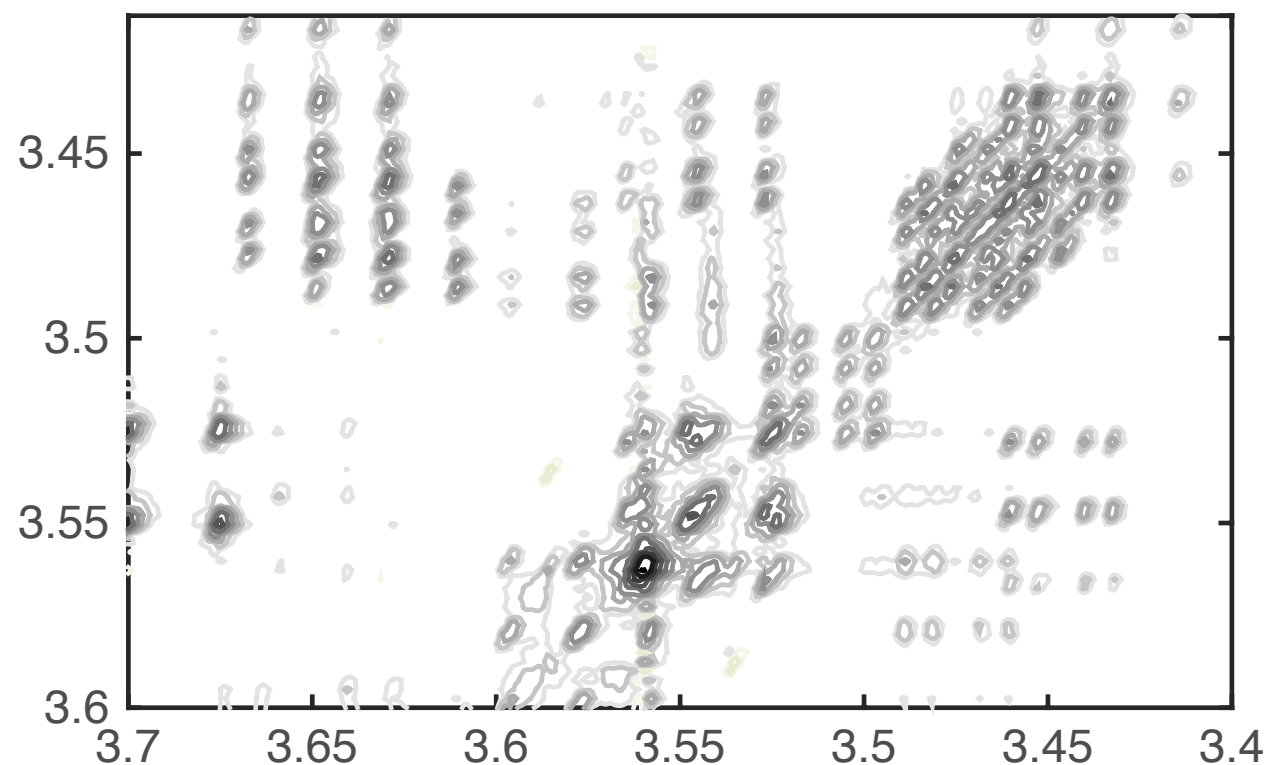
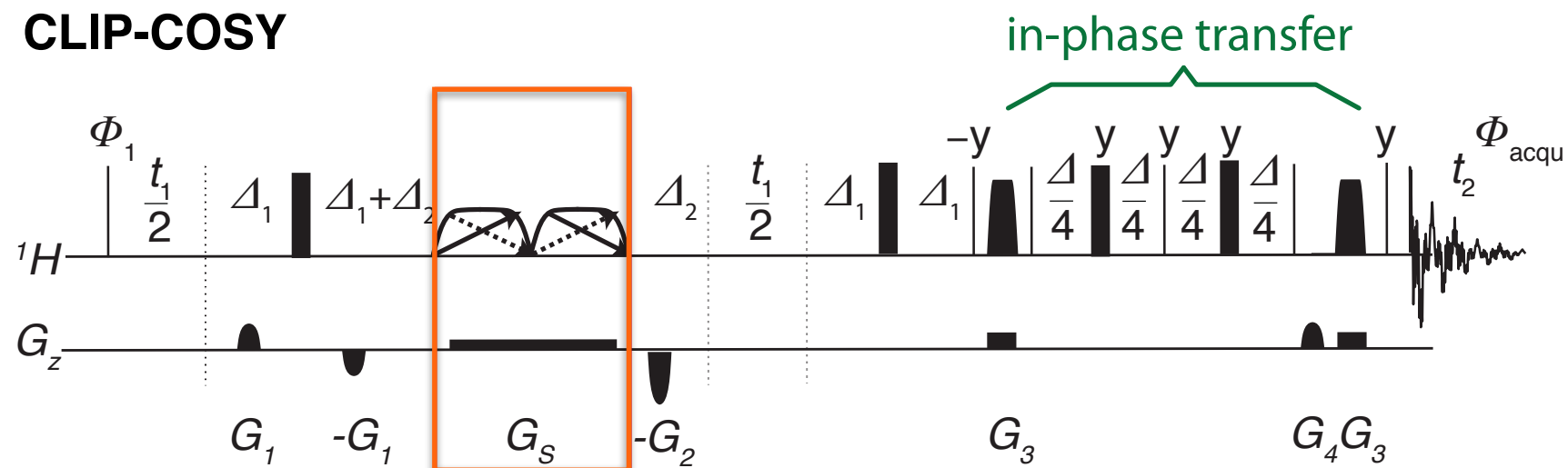
Damien Jeannerat

*Departement of Organic Chemistry,
University of Geneva, Switzerland*



Simplified structure by *Homodecoupling*

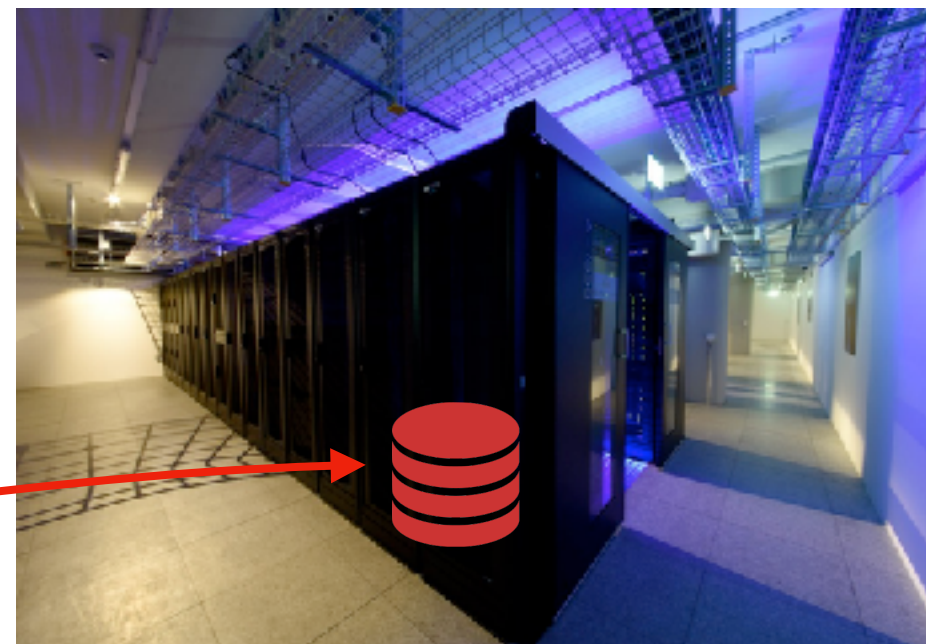
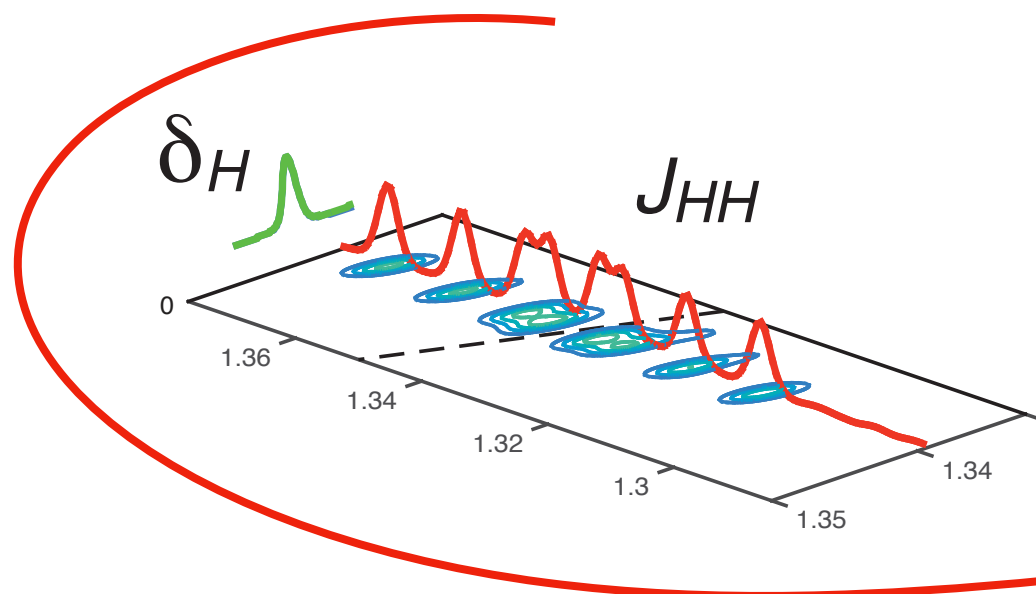
CLIP-COSY



M. R. M. Koos, G. Kummerlowe, L. Kalschnee, C. M. Thiele, B. Luy, Angew. Chem. Int. Ed. 2016, 55.

High-quality spectra

1D ^1H , ^{13}C , DEPT
 2D COSY
 2D HSQC / HMBC ...
 2D J-Resolved / DIAG



High-quality spectra

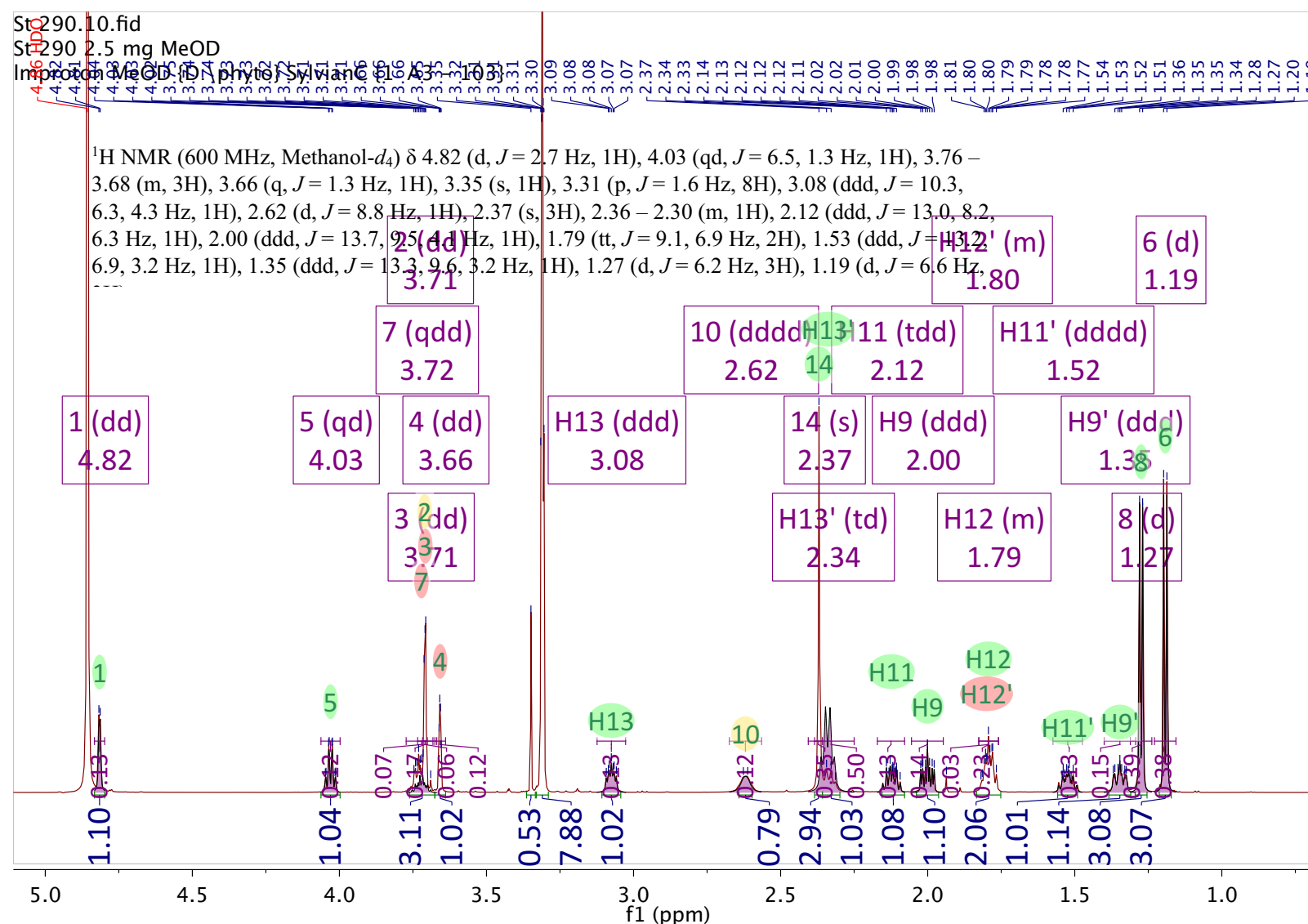
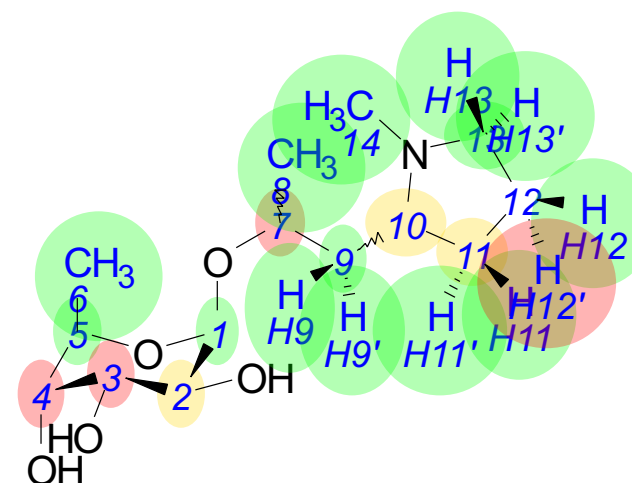
1D ^1H , ^{13}C , DEPT
2D COSY
2D HSQC / HMBC ...
2D J-Resolved / DIAG

Extract information

Chemical shifts
Scalar couplings
2D correlations
...

Share information

Spectrometer
NMR technician
Chemist
Supervisor
Editor
Reviewer
Publisher
Reader / user



High-quality spectra

1D ^1H , ^{13}C , DEPT
2D COSY
2D HSQC / HMBC ...
2D J-Resolved / DIAG

Extract information

Chemical shifts
Scalar couplings
2D correlations
...

Share information

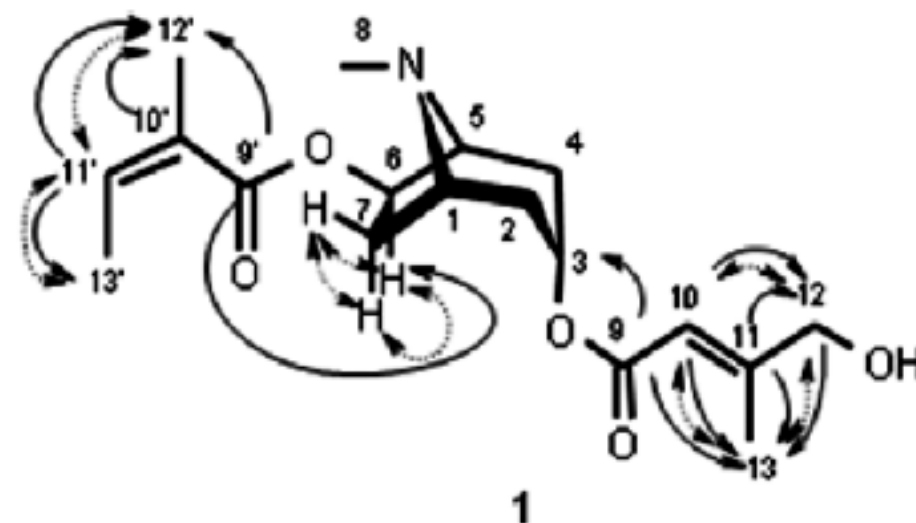
Spectrometer
NMR technician
Chemist
Supervisor
Editor
Reviewer
Publisher
Reader / user

3 α -(E)-4-Hydroxyseneciolyloxy-6 β -angeloyloxytropene (1): colorless oil; $[\alpha]_D^{25}$ -6.25 (MeOH, c 0.04); UV (MeOH) λ_{max} (log ϵ) 220 nm (2.6); **^1H NMR and ^{13}C NMR, see Table 1**; $\text{RI}_{\text{PT}} = 2492.4$; EIMS m/z (rel int) 337 $[\text{M}]^{++}$ (1), 320 (1), 238 (3), 222 (5), 138 (12), 122 (28), 96 (28), 73 (14), 55 (100) ($\text{C}_{18}\text{H}_{28}\text{O}_4$).

Table 1. NMR Spectroscopic Data (500 MHz, CD_3OD , δ in ppm) for Compounds 1–3 Obtained Using Capillary NMR

	1		2		3	
position	δ_{C}^a	δ_{H}	δ_{C}^a	δ_{H}	δ_{C}^a	δ_{H}
1	61.4 CH	3.76 s	61.9	3.92 s	60.0 ^c	4.02 s
2	33.7 CH _{endo}	2.34 s	33.4	2.39 br s		2.43 br s
	CH _{exo}	1.94 s		2.02 s		2.29–2.31 br s
3	64.3 CH	5.08, s	63.8	5.10 s	62.5 ^a	5.14 s
4	32.3 CH _{endo}					
	CH _{exo}					
5	66.9 CH					
6	76.5 CH					
7	34.2 CH _{endo}					
	CH _{exo}					
8	39.0 CH ₃					
9	165.5 ^b qC					
10	112.4 CH					
11	160.5 ^c qC					
12	65.9 CH ₂					
	OH					
13	14.6 CH ₃					
9'	167.4 ^c qC					
10'	127.3 ^c qC					
11'	139.0 CH					
12'	19.4 CH ₃					
13'	26.1 CH ₃	2.01 s	26.3	1.95 s	26.2 ^c	1.95 s

^a Based on HSQC. ^b Based on standard ^{13}C NMR spectra measured in CDCl_3 for the isomer mixture (isomer subfraction). ^c Based on HMBC.



Jeannerat, D., Human- and computer-accessible 2D correlation data for a more reliable structure determination of organic compounds. Future roles of researchers, software developers, spectrometer managers, journal editors, reviewers, publisher and database managers toward artificial-intelligence analysis of NMR spectra. *Magn. Reson. Chem.* **2017**, 55 (1), 7-14.

Academic partners:

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Christoph Steinbeck⁶, Antony J. Williams^{9,&}, Craig Butts¹⁰, Bozhana Mikhova^{11,*}, Damien Jeannerat^{4,§}

Industrial partners:

Wolfgang Robien

Kessler Pavel[#], Fabrice Moriaud,[#] Mikhail Elyashberg[%], Dimitris Argyropoulos[%], Manuel Pérez^{&§}

Publishing partner:

Patrick Giraudeau^{7,§}, Roberto R. Gil^{8,@}, Gary Martin^{‡,@}, Paul Trevorrow[£]



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Magnetic Resonance in Chemistry, Wiley, Chichester, UK

[@] Coeditors in Chief

^{*} Member of the *Editorial board*

[§] Member of the *Associate board*

[&] Member of the *Advisory board*

[£] Executive journal editor

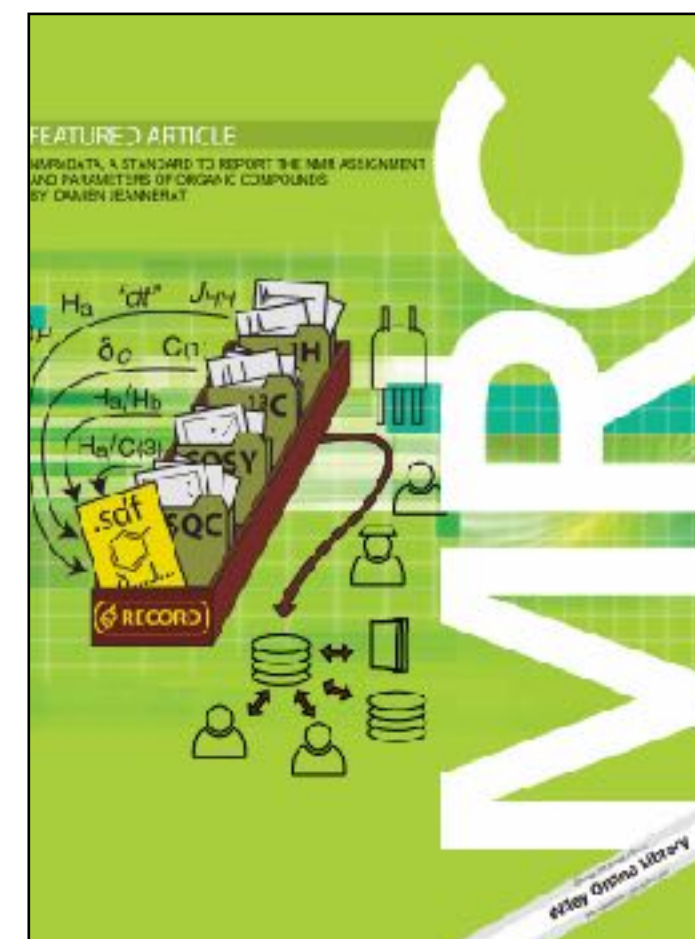
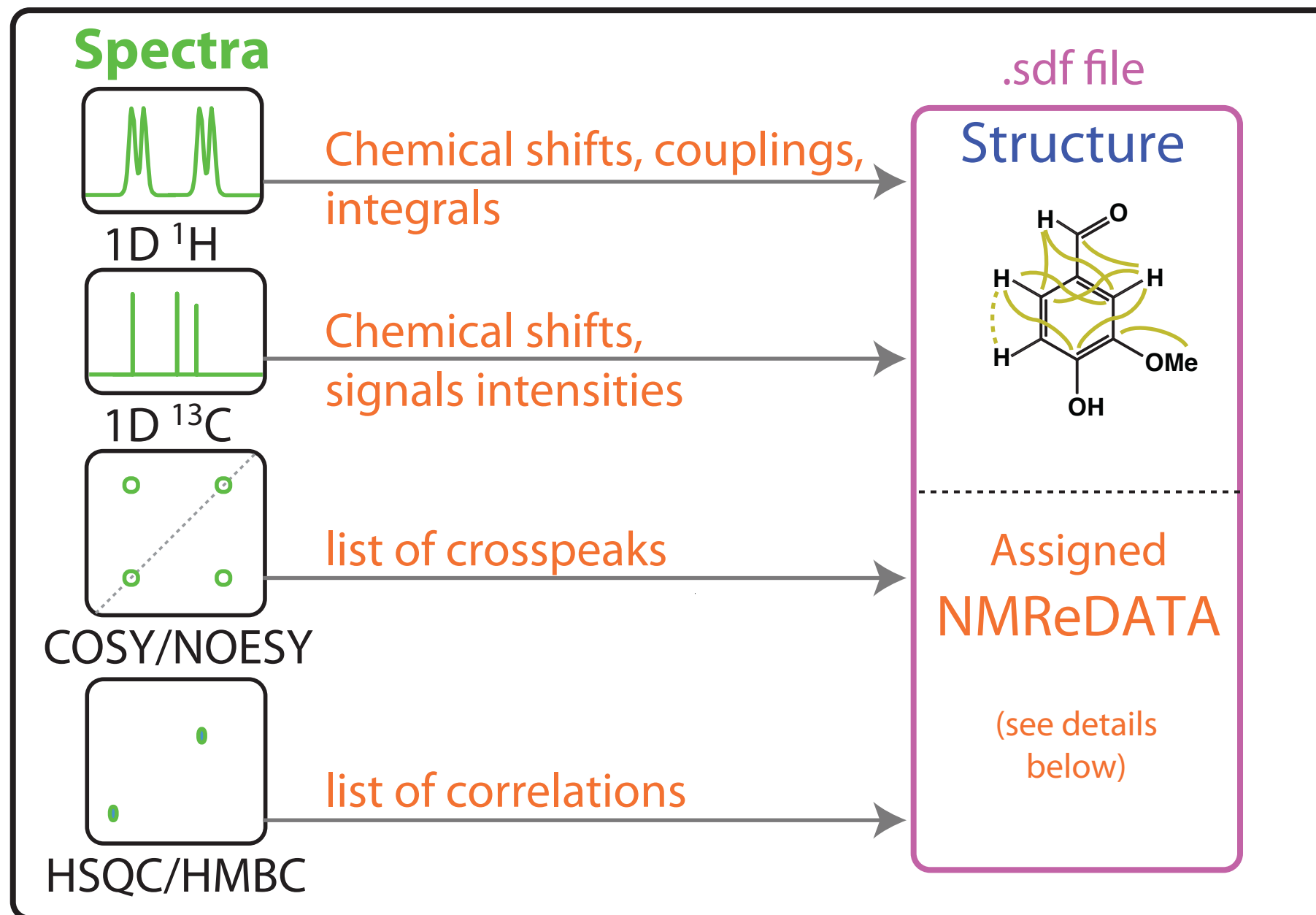


POSTER 376

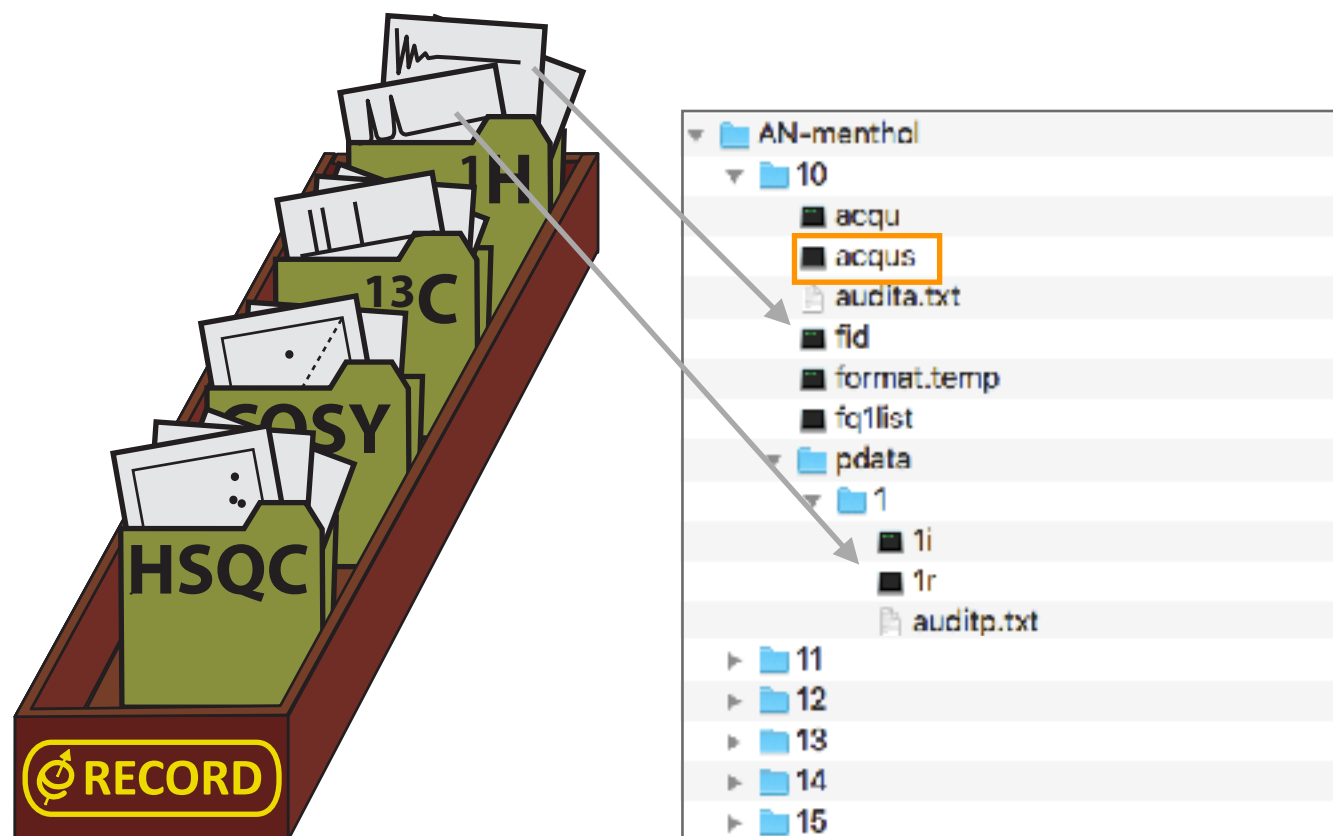
Follow the progress of the initiative on
www.nmredata.org

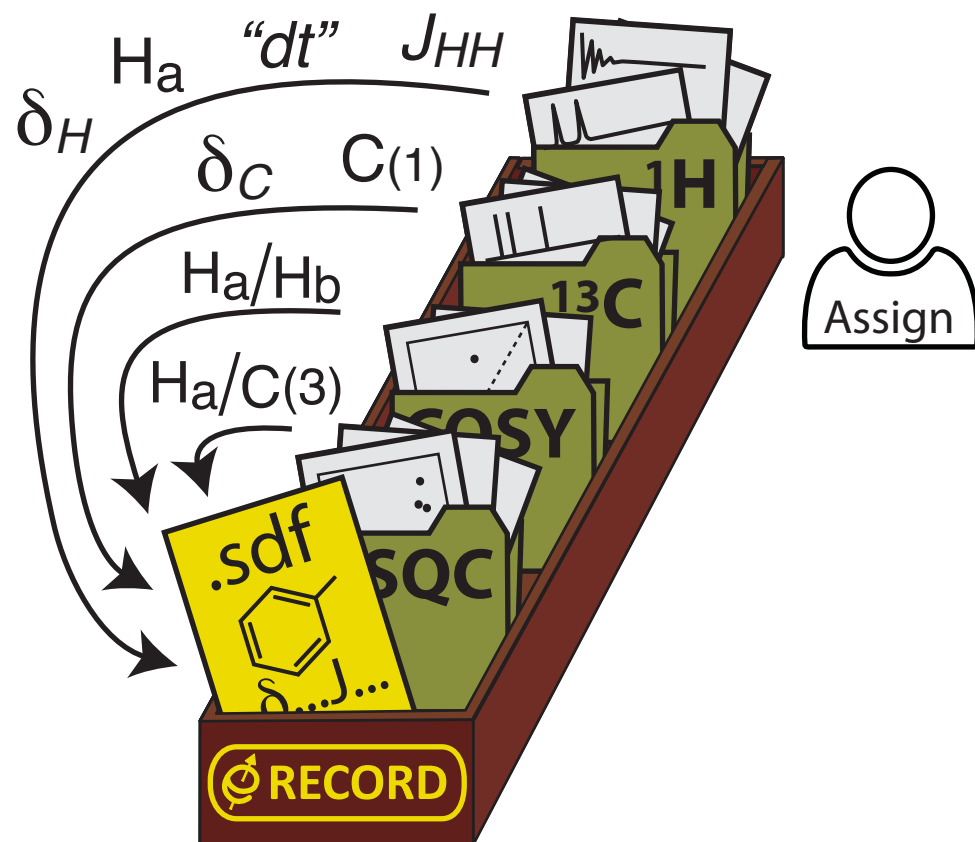


NMR record



1. Pupier, M.; Nuzillard, J.-M.; Wist, J.; Schlörer, N. E.; Kuhn, S.; Erdelyi, M.; Steinbeck, C.; Williams, A. J.; Butts, C.; Claridge, T. D. W.; Mikhova, B.; Robien, W.; Dashti, H.; Eghbalnia, H. R.; Farès, C.; Adam, C.; Kessler, P.; Moriaud, F.; Elyashberg, M.; Argyropoulos, D.; Pérez, M.; Giraudeau, P.; Gil, R. R.; Trevorow, P.; Jeannerat, D., NMReDATA, a standard to report the NMR assignment and parameters of organic compounds. *Magn. Reson. in Chem.* **2018**. DOI: 10.1002/mrc.4737





benzo(a)pyrene
demo of .sdf file containing NMReDATA

test_generationV1 1

20 24 0 0 0 0 0 0 0 0 0999 V2000

-1.6583	2.2334	0.0000	C	0	0	0	0	0	0	0	0	0	0	0	0	0
-0.6633	3.1556	0.0000	C	0	0	0	0	0	0	0	0	0	0	0	0	0
-1.3753	0.8191	0.0000	C	0	0	0	0	0	0	0	0	0	0	0	0	0
0.7234	2.7603	0.0000	C	0	0	0	0	0	0	0	0	0	0	0	0	0
0.0000	0.3848	0.0000	C	0	0	0	0	0	0	0	0	0	0	0	0	0

...

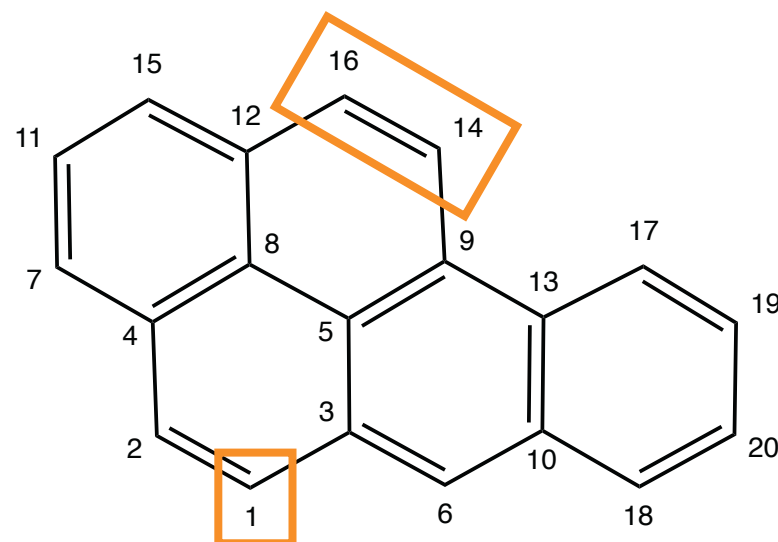
2.6960	-0.4398	0.0000	C	0	0	0	0	0	0	0	0	0	0	0	0	0
-0.5264	-3.3746	0.0000	C	0	0	0	0	0	0	0	0	0	0	0	0	0
-3.1687	-2.4653	0.0000	C	0	0	0	0	0	0	0	0	0	0	0	0	0
-1.5724	-4.2723	0.0000	C	0	0	0	0	0	0	0	0	0	0	0	0	0
-2.9104	-3.8155	0.0000	C	0	0	0	0	0	0	0	0	0	0	0	0	0

1	2	2	0	0	0
1	3	1	0	0	0
2	4	1	0	0	0
3	5	1	0	0	0

...

13	17	1	0	0	0
14	16	2	0	0	0
17	19	2	0	0	0
18	20	2	0	0	0
19	20	1	0	0	0

M END



.mol

.sdf

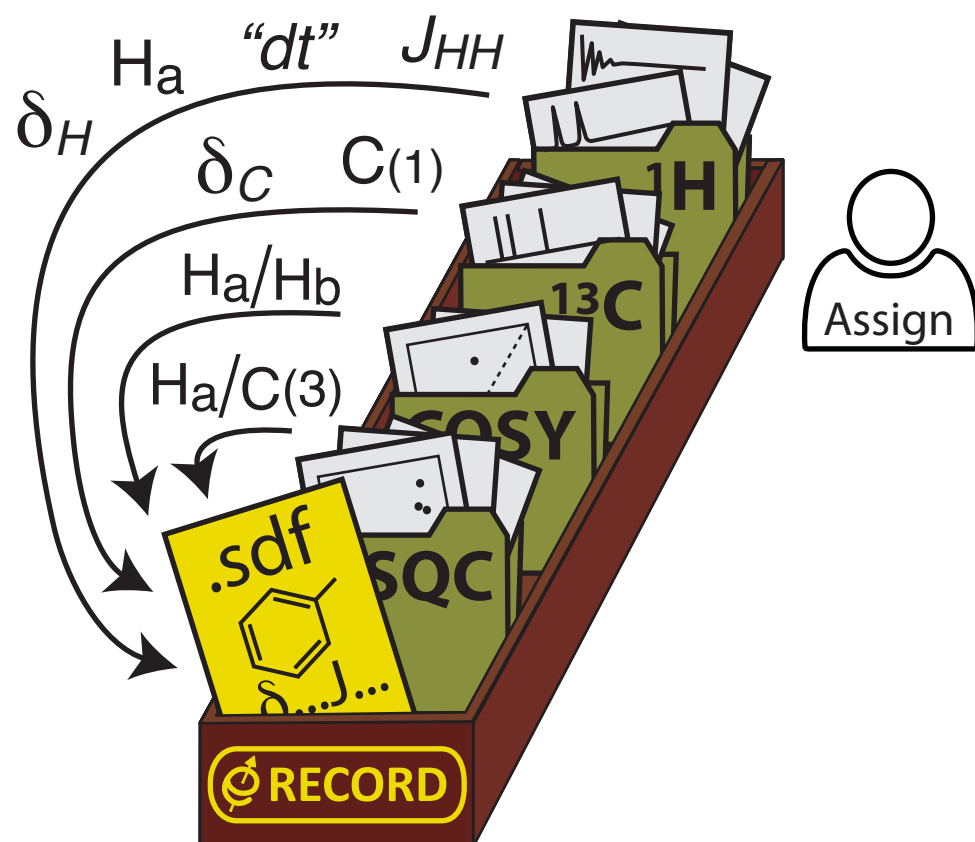
Structure
Data
Format

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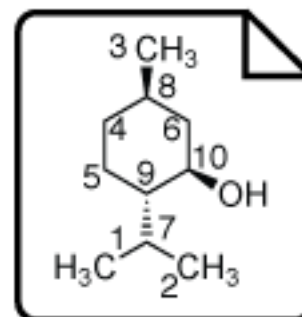
cdcl3

> <NMREDATA_TEMPERATURE>

298.15



Assigned NMReDATA [compound1.nmredata.sdf](#)



NMReDATA_1D_13C

NMReDATA_1D_1H

1.1301, S=ddd, L=H9ax, E=42.3746, J=12.76, 9.90, 3.05, 2.72
 3.4302, S=dddd, L=H10ax, E=28.9715, J=10.90, 9.90, 4.80, 4.50
 1.4444, S=ddqdd, L=H8ax, E=27.4764, J=12.00, 12.00, 6.58, 4.00, 3.00
 ...

NMReDATA_2D_13C_NJ_1H

NMReDATA_2D_13C_1J_1H

C3/Me3
 C4/H4ax
 C4/H4eq
 ...


```
> <NMREDATA_1D_1H>
```

```
Larmor=500.133088507\
```

```
Pulseprogram=zg30
```

```
Spectrum_Location=file:AN-menthol/10/pdata/1/\
```

```
1.1301, S=dddd, L=H3, E=42.3746, J=12.76,9.90,3.05,2.72\
```

```
3.4302, S=dddd, L=H4, E=28.9715, J=10.90,9.90,4.80,4.30\
```

```
1.4444, S=ddtdd, L=H6, E=27.4764, J=12.00,12.00,6.58,4.00,3.00\
```

```
2.1895, S=ddd, L=H9, E=42.6060, J=7.02,6.90,2.72\
```

```
...
```

```
> <NMREDATA_1D_13C>
```

```
Larmor=125.770363831\
```

```
Pulseprogram=dept135 ;optional in V1\
```

```
Spectrum_Location=file:AN-menthol/12/pdata/1/\
```

```
...
```

```
> <NMREDATA_2D_1H_NJ_1H>
```

```
Larmor=500.13\
```

```
CorrType=COSY\
```

```
Pulseprogram=cosygpmfph2
```

```
Spectrum_Location=file:dj_menthol/21/pdata/1/\
```

```
H4/H3\
```

```
H9/H3\
```

```
H3/H4\
```

```
H12/H4\
```

```
H5ax/H4\
```

```
H5eq/H4\
```

```
H7/H6\
```

```
> <NMREDATA_1D_1H>
```

```
Larmor=500.133088507\
```

```
Pulseprogram=zg30
```

```
Spectrum_Location=file:AN-menthol/10/pdata/1/\
```

```
1.1301, S=dddd, L=H3, E=42.3746, J=12.76,9.90,3.05,2.72\
```

```
3.4302, S=dddd, L=H4, E=28.9715, J=10.90,9.90,4.80,4.30\
```

```
1.4444, S=ddtdd, L=H6, E=27.4764, J=12.00,12.00,6.58,4.00,3.00\
```

```
2.1895, S=ddd, L=H9, E=42.6060, J=7.02,6.90,2.72\
```

```
...
```

```
> <NMREDATA_1D_13C>
```

```
Larmor=125.770363831\
```

```
Pulseprogram=dept135 ;optional in V1\
```

```
Spectrum_Location=file:AN-menthol/12/pdata/1/\
```

```
...
```

```
> <NMREDATA_2D_1H_NJ_1H>
```

```
Larmor=500.13\
```

```
CorrType=COSY\
```

```
Pulseprogram=cosygpmfph2
```

```
Spectrum_Location=file:dj_menthol/21/pdata/1/\
```

```
H4/H3\
```

```
H9/H3\
```

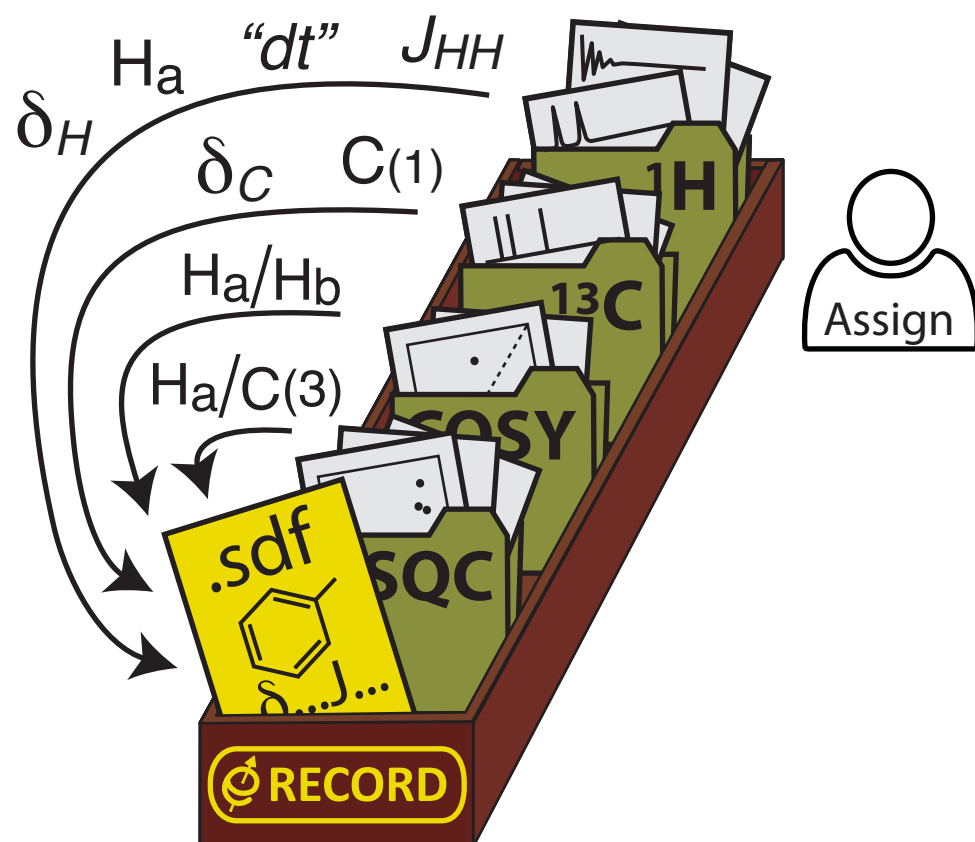
```
H3/H4\
```

```
H12/H4\
```

```
H5ax/H4\
```

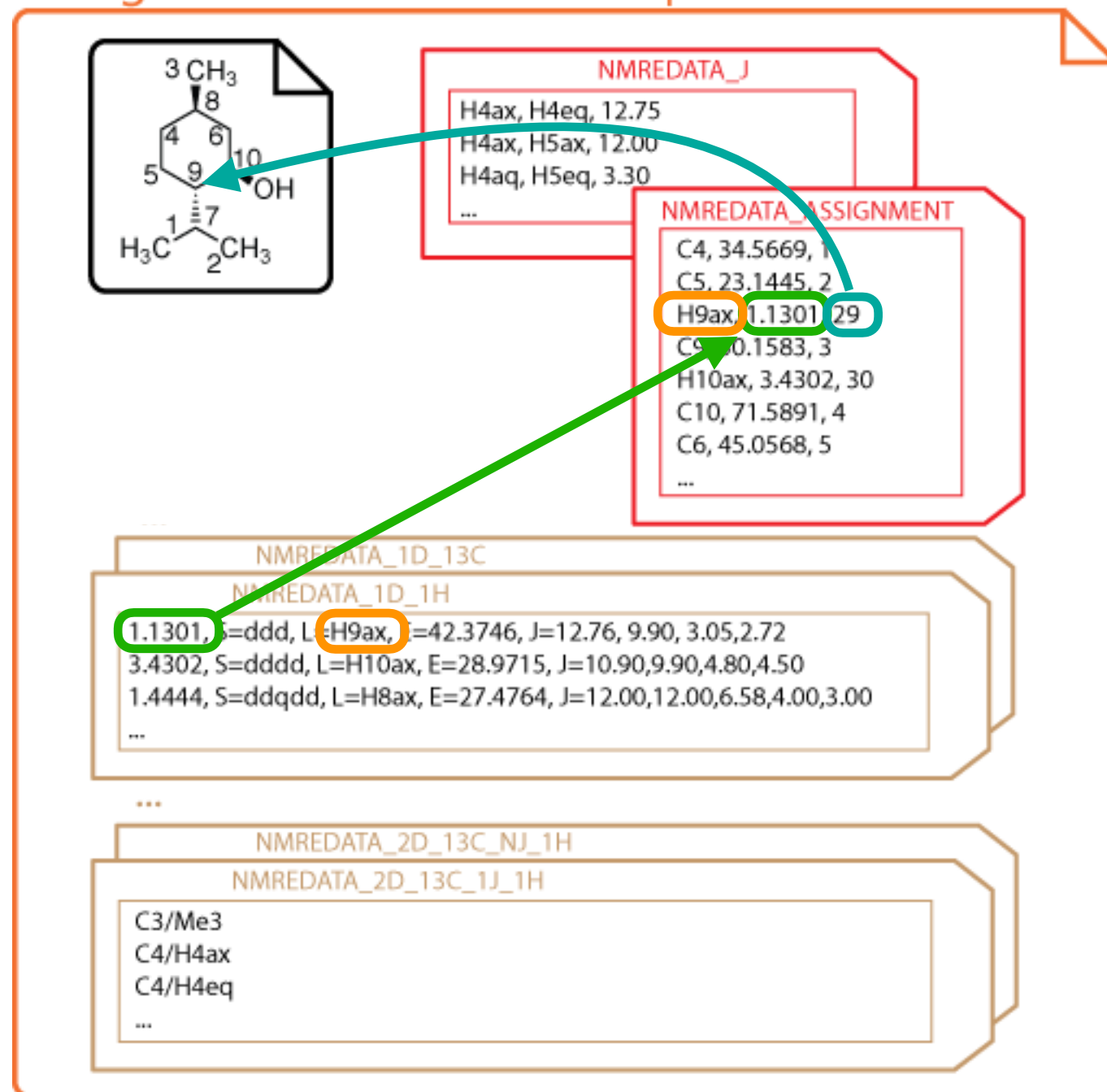
```
H5eq/H4\
```

```
H7/H6\
```



Desired redundance and complementarity

Assigned NMReDATA compound1.nmredata.sdf



Label / δ / Atom number

```
> <NMREDATA_ASSIGNMENT>
```

```
1, 34.5669, 1\
```

```
2, 23.1445, 2\
```

```
H3, 1.1301, H3\
```

```
3, 50.1583, 3\
```

When H atoms are implicit add "H"...

```
H4, 3.4302, H4\
```

```
4, 71.5891, 4\
```

```
5, 45.0568, 5\
```

```
H6, 1.4444, H6\
```

```
6, 31.6232, 6\
```

```
H7, 0.9331, H7\
```

```
7, 22.2293, 7\
```

```
H12, 1.3536, H8\
```

```
H9, 2.1895, H9\
```

```
9, 25.8422, 9\
```

```
H10, 0.8311, H10\
```

```
10, 16.1017, 10\
```

```
H11, 0.9493, H11\
```

```
11, 20.9880, 11\
```

```
H1eq, 1.6822, 12\
```

```
H1ax, 0.8630, 13\
```

```
H2ax, 0.9933, 14\
```

```
H2eq, 1.6293, 15\
```

```
H5ax, 0.9535, 16\
```

```
H5eq, 1.9844, 17\
```


Label / δ / Atom number

> <NMREDATA_ASSIGNMENT>

```
1, 34.5669, 1\  
2, 23.1445, 2\  
H3, 1.1301, H3\  
3, 50.1583, 3\  
H4, 3.4302, H4\  
4, 71.5891, 4\  
5, 45.0568, 5\  
H6, 1.4444, H6\  
6, 31.6232, 6\  
H7, 0.9331, H7\  
7, 22.2293, 7\  
H12, 1.3536, H8\  
H9, 2.1895, H9\  
9, 25.8422, 9\  
H10, 0.8311, H10\  
10, 16.1017, 10\  
H11, 0.9493, H11\  
11, 20.9880, 11\  
H1eq, 1.6822, 12\  
H1ax, 0.8630, 13\  
H2ax, 0.9933, 14\  
H2eq, 1.6293, 15\  
H5ax, 0.9535, 16\  
H5eq, 1.9844, 17\  

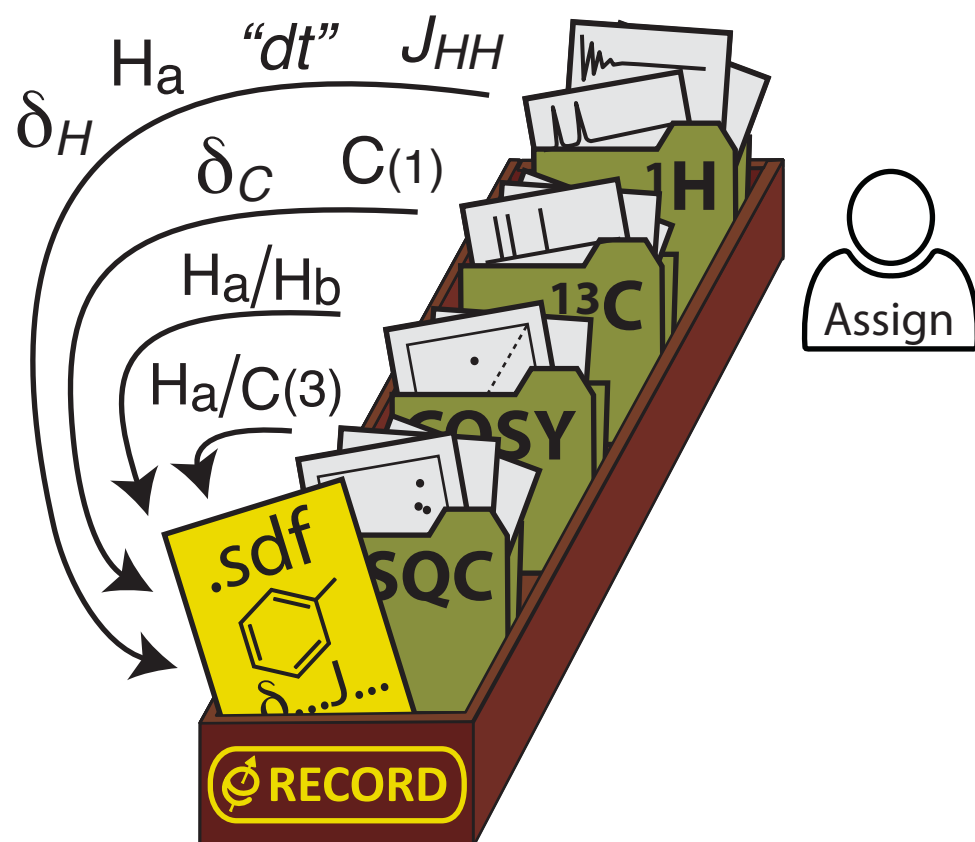
```

> <NMREDATA_J>

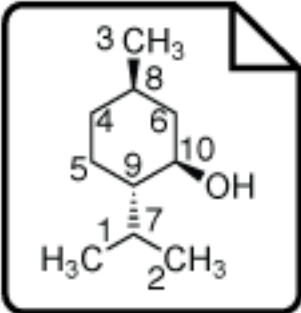
optional

```
H3, H2ax, 12.8\  
H3, H2eq, 3.0\  
H3, H4, 9.9\  
H3, H9, 2.7\  
H4, H12, 4.8\  
H4, H5ax, 10.9\  
H4, H5eq, 4.5\  
H6, H7, 6.6\  
H6, H1eq, 3.0\  
H6, H1ax, 12.0\  
H6, H5ax, 12.0\  
H6, H5eq, 4.0\  
H9, H10, 7.0\  
H9, H11, 7.1\  
H1eq, H1ax, 12.8\  
H1eq, H2ax, 3.3\  
H1eq, H2eq, 3.2\  
H1eq, H5eq, 2.2\  
H1ax, H2ax, 12.0\  
H1ax, H2eq, 3.3\  
H2ax, H2eq, 13.0\  
H5ax, H5eq, 12.1\  

```



Assigned NMReDATA compound1.nmredata.sdf



...

NMREDATA_J

H4ax, H4eq, 12.75
H4ax, H5ax, 12.00
H4aq, H5eq, 3.30
...

NMREDATA_ASSIGNMENT

C4, 34.5669, 1
C5, 23.1445, 2
H9ax, 1.1301, 29
C9, 50.1583, 3
H10ax, 3.4302, 30
C10, 71.5891, 4
C6, 45.0568, 5
...

NMREDATA_TEMPERATURE

NMREDATA_SOLVENT

CDCL3

...

NMREDATA_1D_13C

NMREDATA_1D_1H

1.1301, S=ddd, L=H9ax, E=42.3746, J=12.76, 9.90, 3.05, 2.72
3.4302, S=dddd, L=H10ax, E=28.9715, J=10.90, 9.90, 4.80, 4.50
1.4444, S=ddqdd, L=H8ax, E=27.4764, J=12.00, 12.00, 6.58, 4.00, 3.00
...

...

NMREDATA_2D_13C_NJ_1H

NMREDATA_2D_13C_1J_1H

C3/Me3
C4/H4ax
C4/H4eq
...

```
M  ZZC  13  1ax  
M  ZZC  14  2ax  
M  ZZC  15  2eq  
M  ZZC  16  5ax  
M  ZZC  17  5eq  
M  END
```

```
> <NMREDATA_VERSION>
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```
1.1\  

```

```
> <NMREDATA_ID>
```

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> <NMREDATA_SOLVENT>
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```
CDC13\  

```

```
> <NMREDATA_LEVEL>
```

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0\  

```

```
> <NMREDATA_ASSIGNMENT>
```

```
1, 34.5669, 1\  

```

```
2, 23.1445, 2\  

```

```
H3, 1.1301, H3\  

```

```
3, 50.1583, 3\  

```

```
H4, 3.4302, H4\  

```

```
4, 71.5891, 4\  

```

```
5, 45.0568, 5\  

```

```
H6, 1.4444, H6\  

```

```
6, 31.6232, 6\  

```

```
H7, 0.9331, H7\  

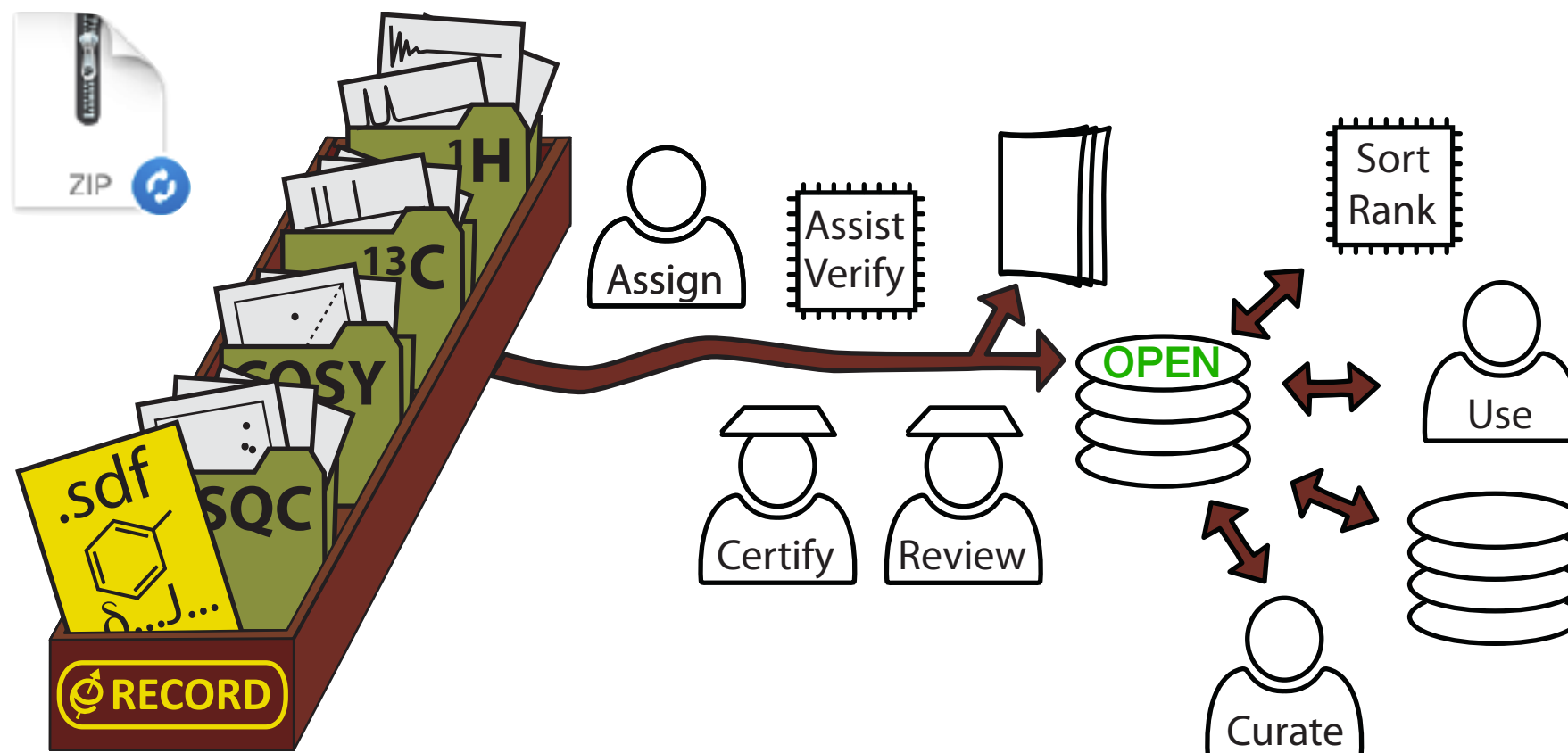
```

```
7, 22.2293, 7\  

```

```
H12, 1.3536, H8\  

```




```
M  ZZC  13  1ax  
M  ZZC  14  2ax  
M  ZZC  15  2eq  
M  ZZC  16  5ax  
M  ZZC  17  5eq  
M  END
```

```
> <NMREDATA_VERSION>
```

```
1.1\
```

```
> <NMREDATA_ID>
```

```
Doi=10.5281/zenodo.1146869
```

```
Record=http://zenodo.org/record/1146869/files/sample1.zip
```

```
> <NMREDATA_SOLVENT>
```

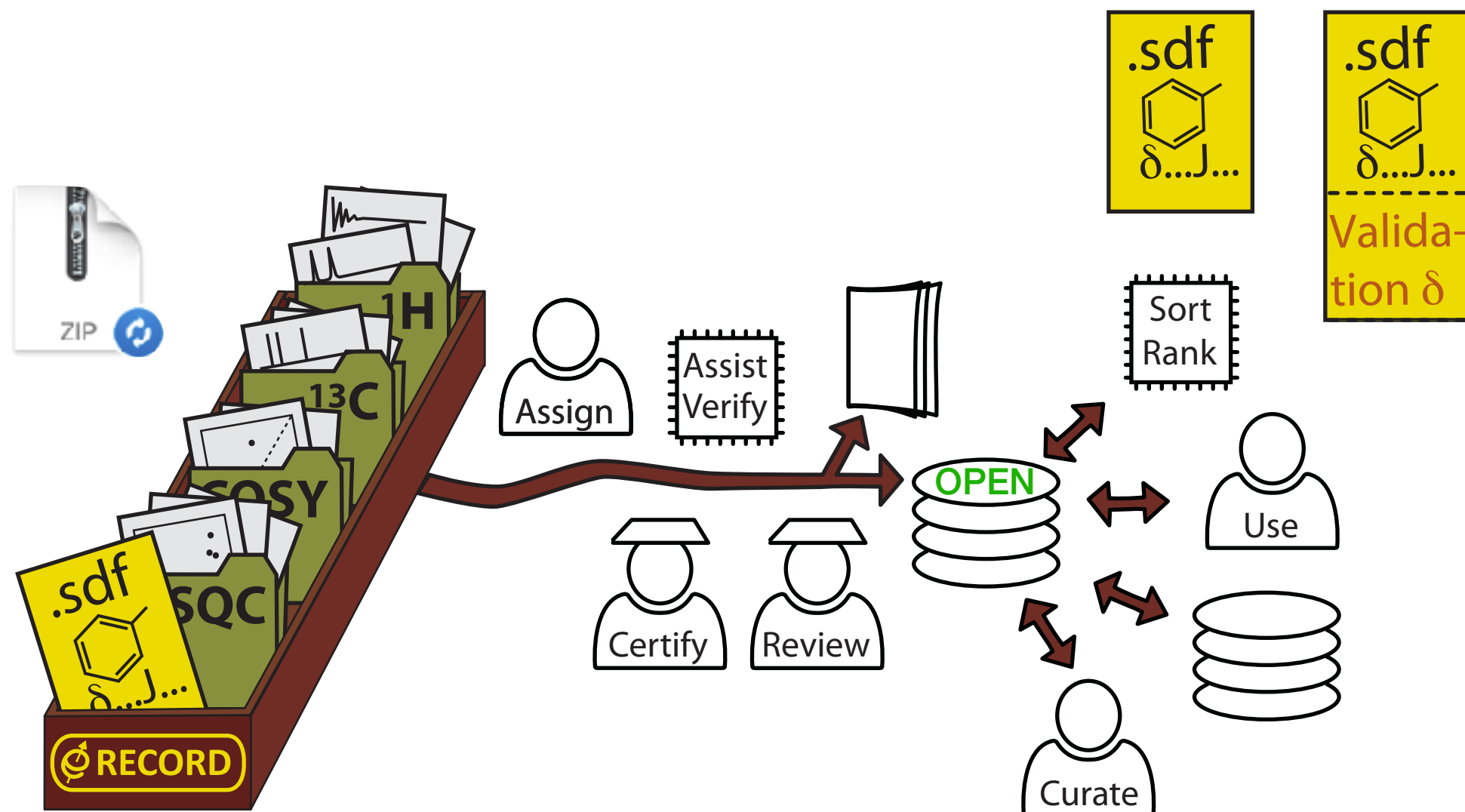
```
CDC13\
```

```
> <NMREDATA_LEVEL>
```

```
0\
```

```
> <NMREDATA_ASSIGNMENT>
```

```
1, 34.5669, 1\  
2, 23.1445, 2\  
H3, 1.1301, H3\  
3, 50.1583, 3\  
H4, 3.4302, H4\  
4, 71.5891, 4\  
5, 45.0568, 5\  
H6, 1.4444, H6\  
6, 31.6232, 6\  
H7, 0.9331, H7\  
7, 22.2293, 7\  
H12, 1.3536, H8\
```

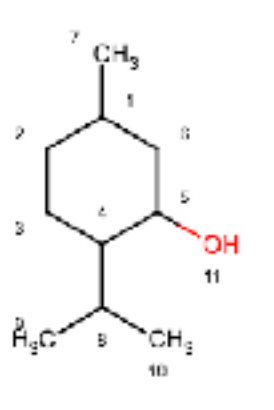


Validation of δ

<http://nmrshiftdb.nmr.uni-koeln.de/>



If you [login](#), your Quick Check results will be available later!



Choose an MNova mol or sd file (MNova 11 or newer) or ChemDraw mol file with labels:
 No file selected.

Choose a Topspin zip file including a CMC-SE result: No file selected.

Choose an NMReDATA file: No file selected.

1D spectra 2D spectra

Atom No.	¹³ C Shift	¹ H Shift
11		Keep unchanged 3.62
1	35.0 Keep unchanged 31.80, diff: 3.20	Keep unchanged 1.44
2	Keep unchanged 34.80, diff: 5.20	Keep unchanged 1.26
3	Keep unchanged 23.35	Keep unchanged 1.30
4	Keep unchanged 50.25	Keep unchanged 1.12
5	Keep unchanged 71.10	Keep unchanged 3.42
6	Keep unchanged 45.35	Keep unchanged 1.47
7	Keep unchanged 22.25	Keep unchanged 0.93
8	Keep unchanged 25.75	Keep unchanged 2.10
9	Keep unchanged + 10.55	Keep unchanged + 0.88
10	Keep unchanged + 10.55	Keep unchanged + 0.88

If your structure has more atoms, transfer to get more fields!
 * = symmetrical atom, if left empty it will be filled in automatically

Submit ¹³C

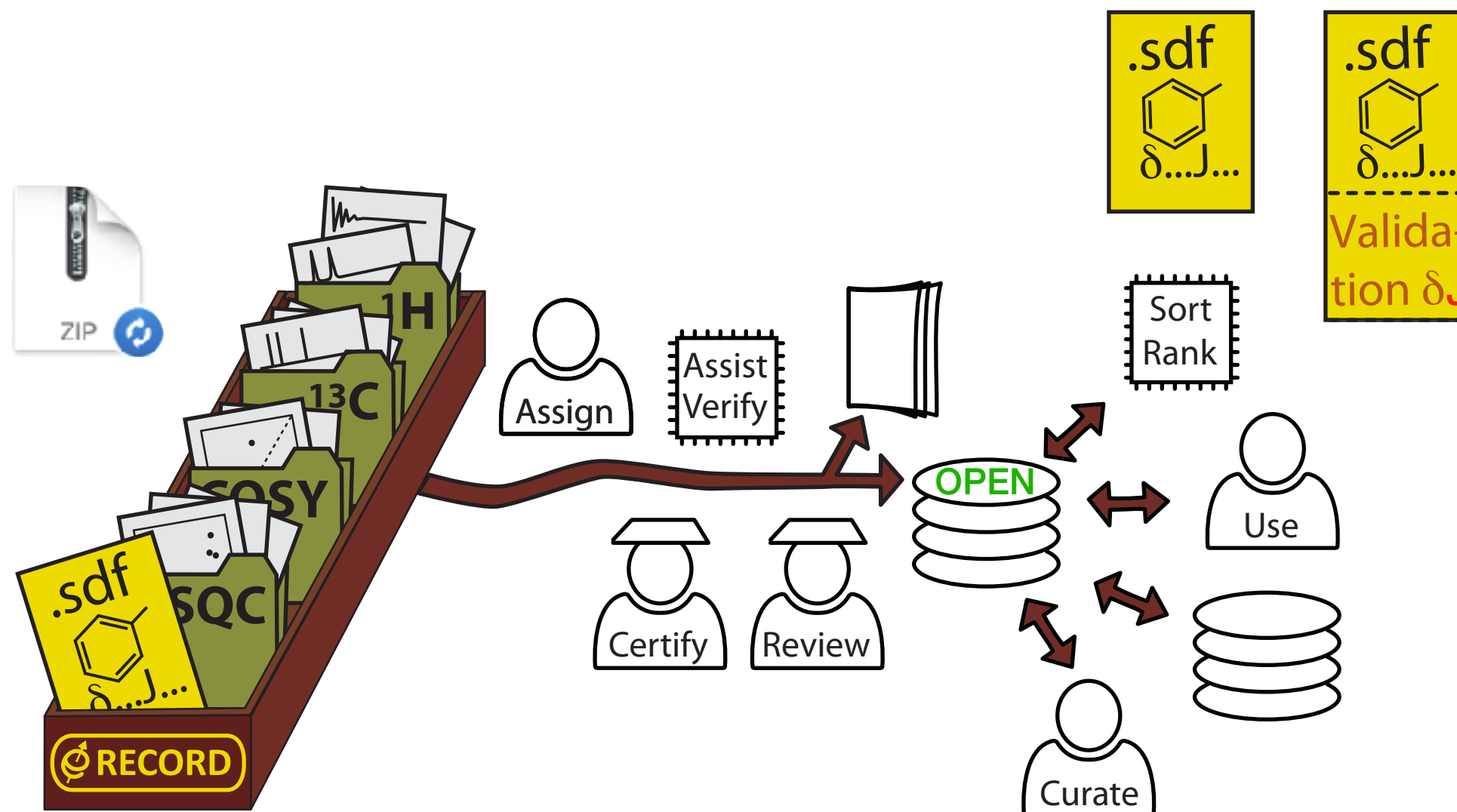
Input list: [Input format](#)

Shifts for auto assignment, format like:
 56.2
 56.25
 64.10
 multiplicity optional

Submit ¹H

Input list: [Input format](#)

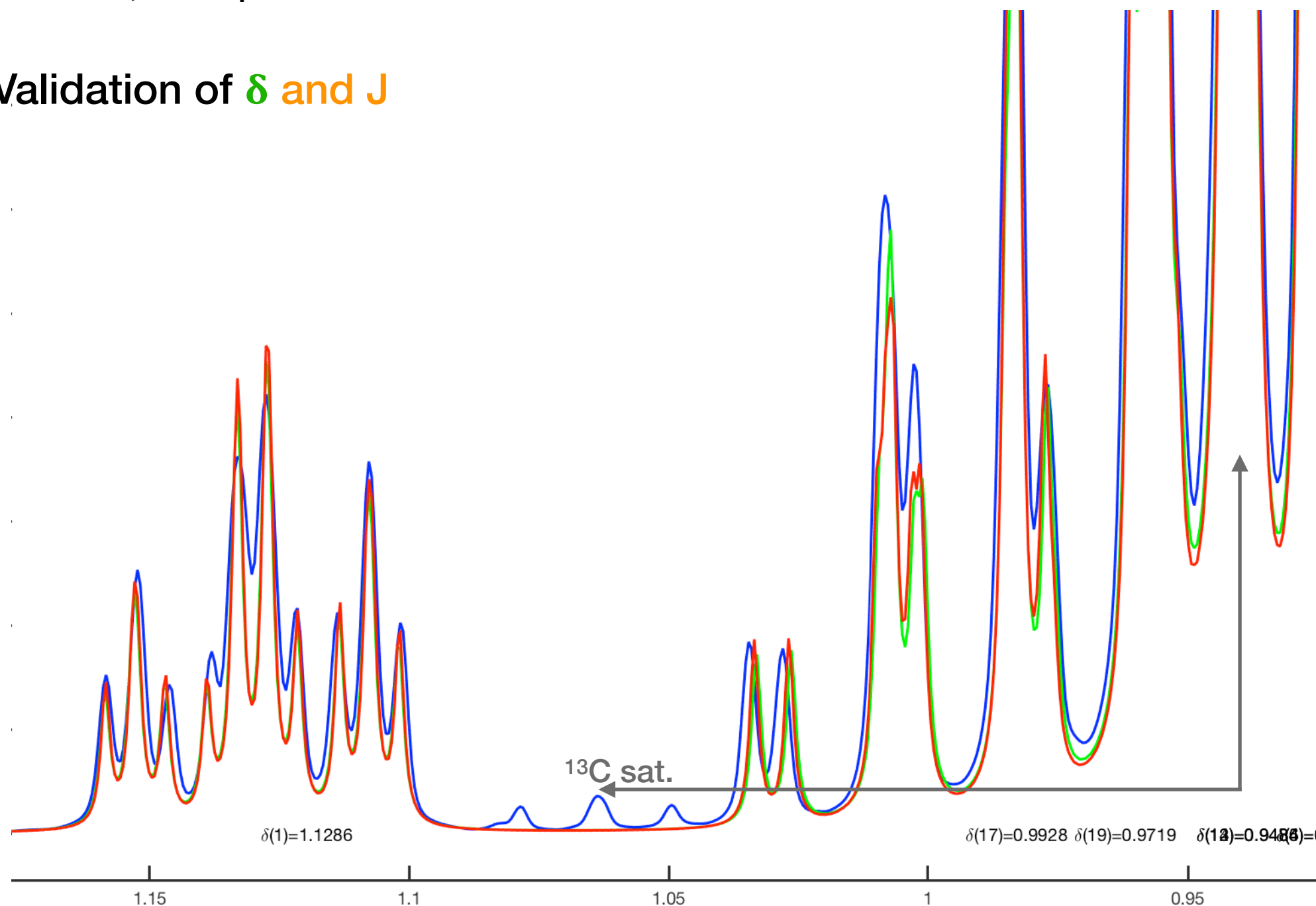
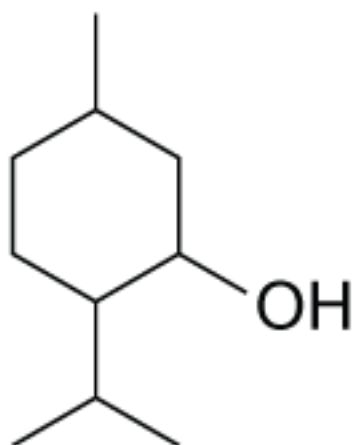
Shifts for auto assignment, format like:
 56.2
 64.1
 Enter diastereotopic pairs like this:
 1.8
 +1.6



Validation of δ and J

... with assignment of J, can reproduce 2nd order effects

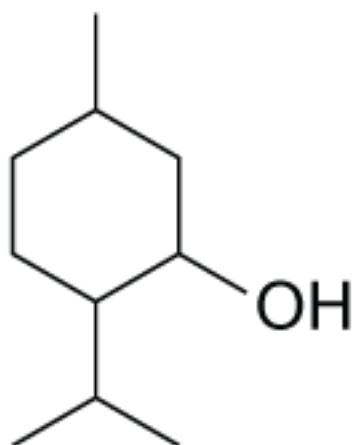
Refinement + Validation of δ and J



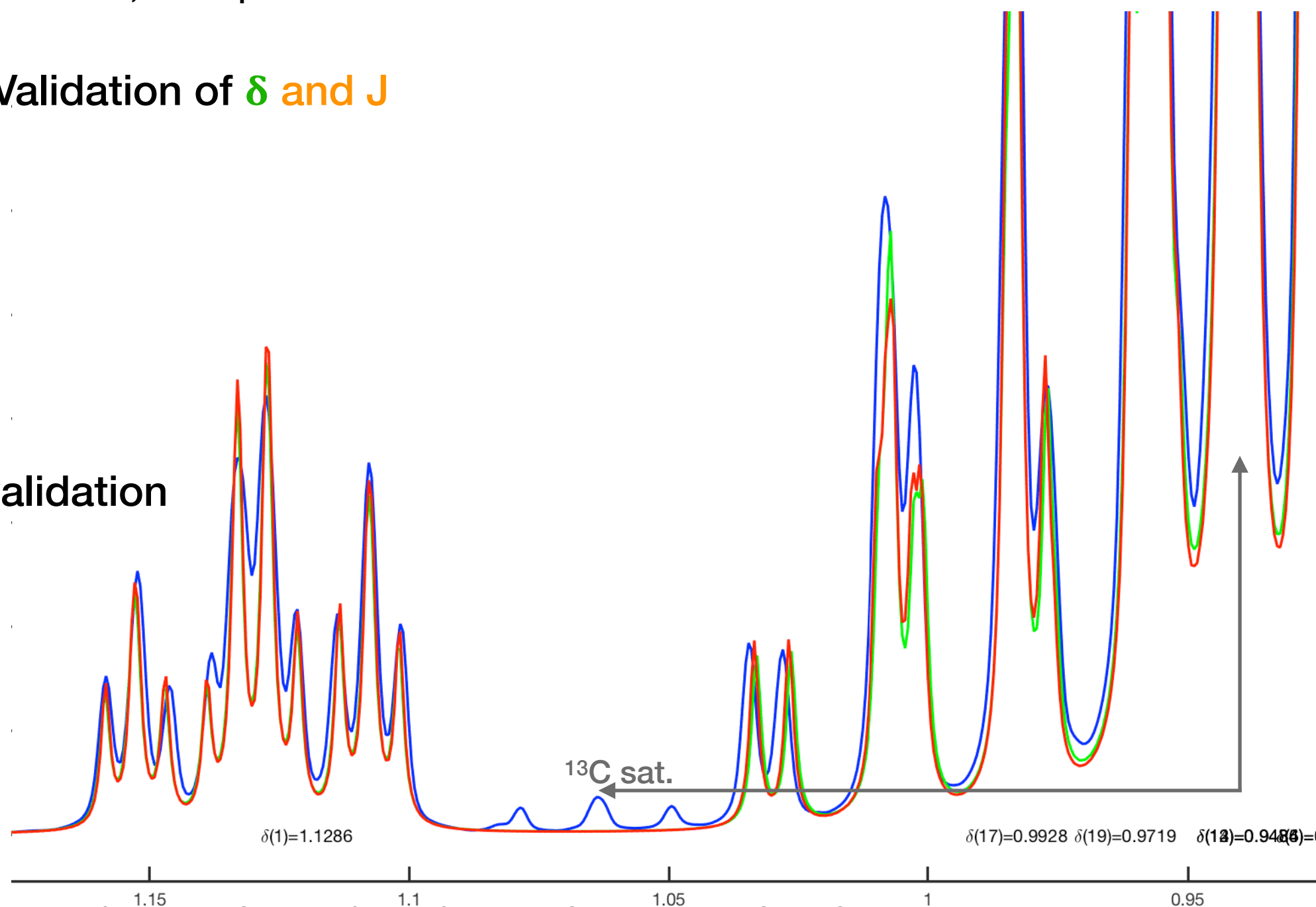
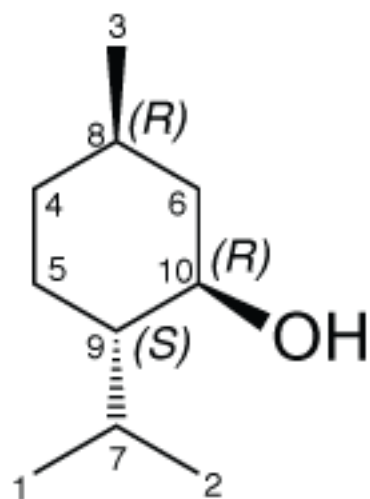
Validation of δ and J

... with assignment of J, can reproduce 2nd order effects

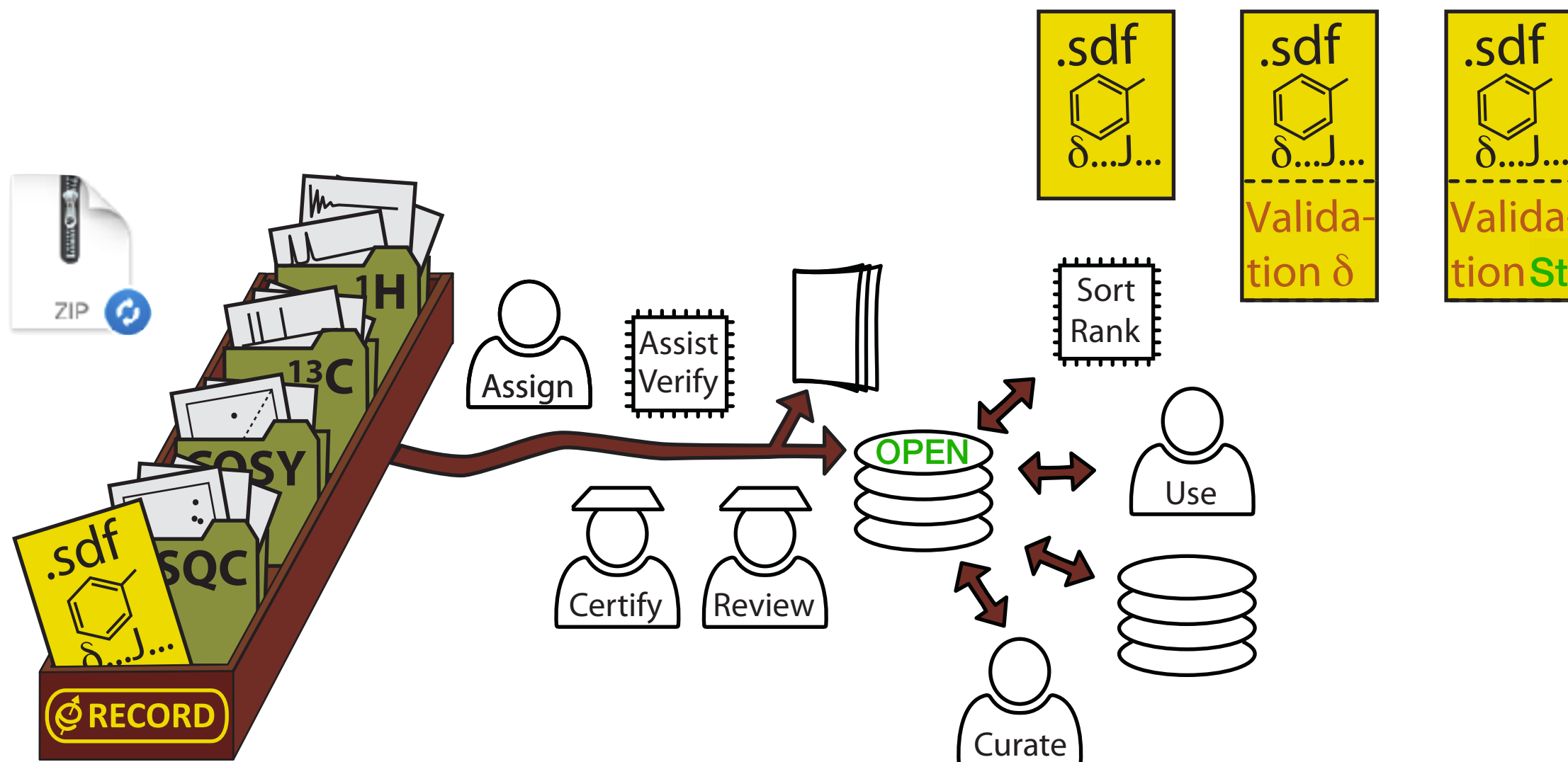
Refinement + Validation of δ and J



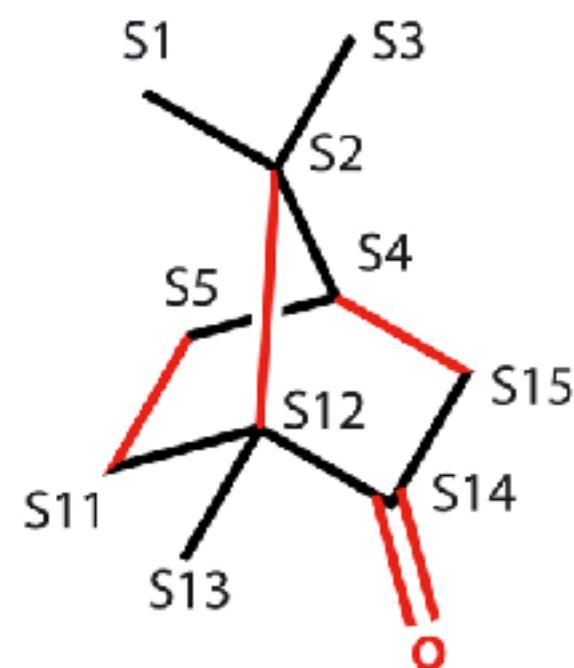
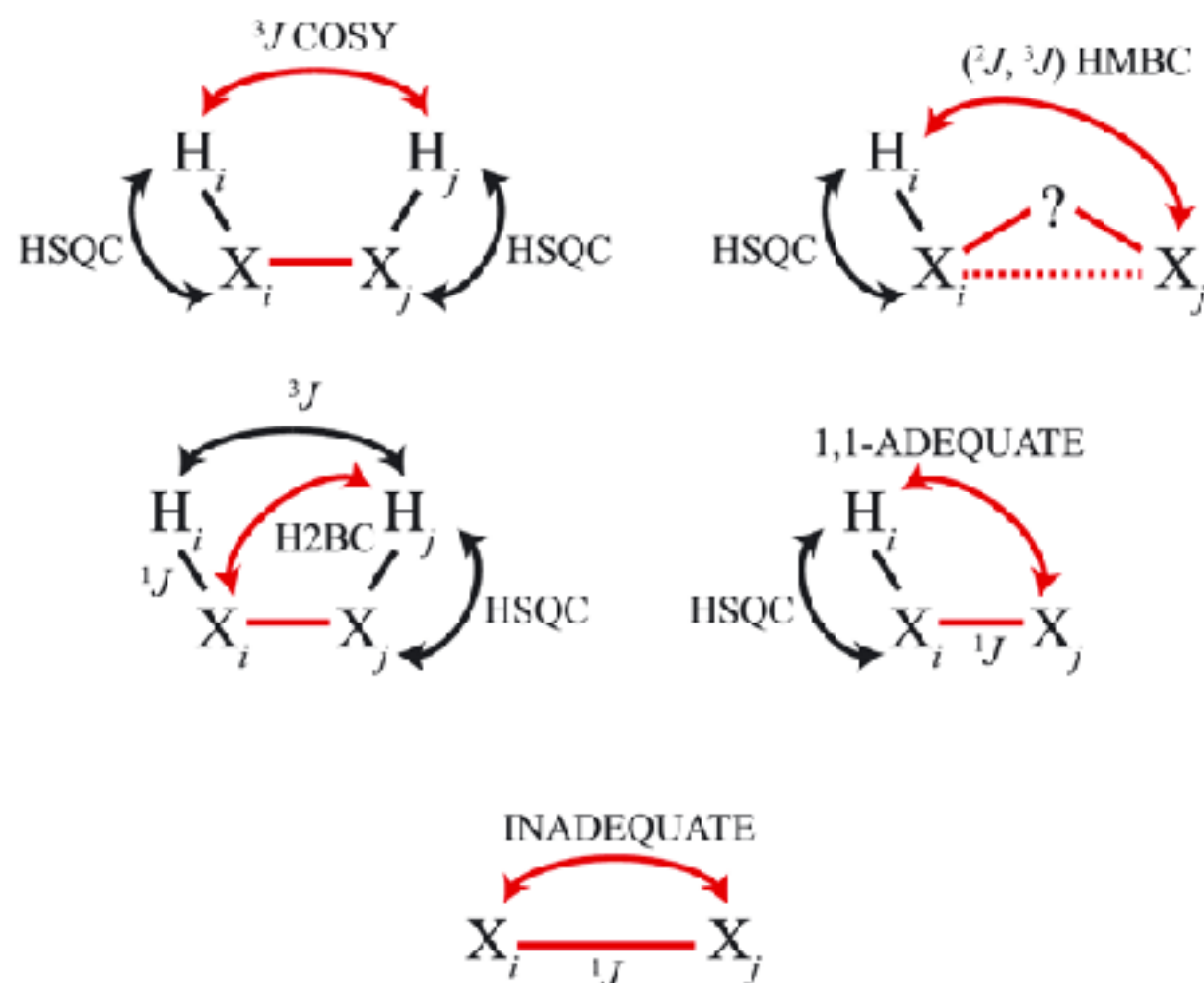
3D structure validation

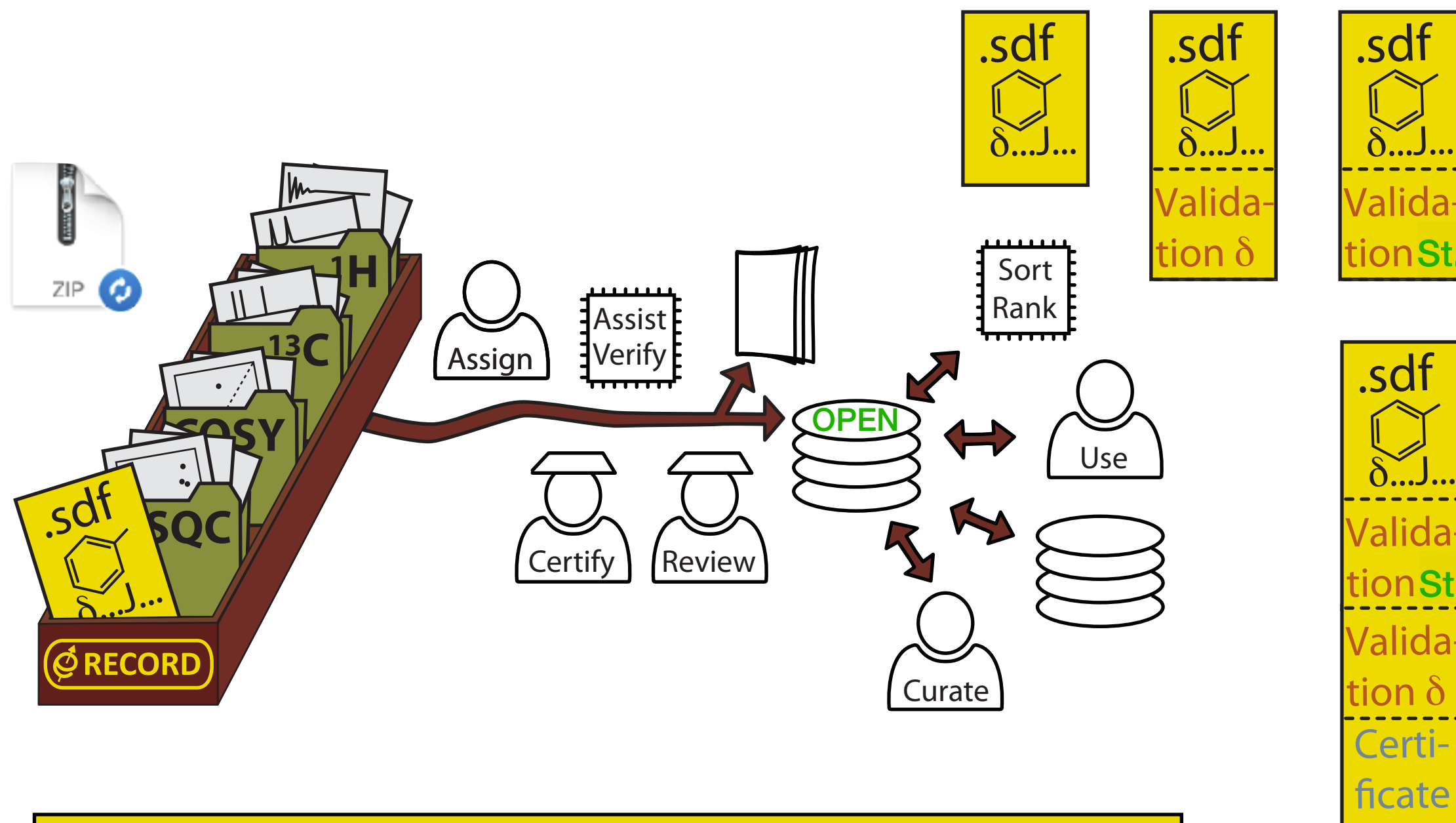


MSpin-JCoupling : Navarro-Vázquez, A.; Santamaría-Fernández, R.; Sardina, F. J., MSpin-JCoupling. A modular program for prediction of scalar couplings and fast implementation of **Karplus** relationships. *Magn. Reson. Chem.* **2017**.



Validation of 2D J correlations using Logic for Structure Determination (LSD) (COSY, HSQC, HMBC, etc.)





NMReDATA

MSeDATA - being considered

REACeDATA - info. about the origin of the sample

NATSOURCEeDATA - Natural products

Compatible software and web service

Working software and tools

Software Platform	Product/tool	Developer	Operating system	Version	NMReDATA	Supported NMReDATA versions	NMR Record	Examples of output
nmrshiftdb2	QuickCheck	Stefan Kuhn	Web browser	4.1.12	generate/write	1.0 & 1.1	generate	link
Javatools GUI and library	see github	Stefan Kuhn	Any Java-enabled	Pre-alpha	generate/write /validate	1.0 & 1.1	generate	
ACD/Spectrus	Spectrus Processor and above	ACD/Labs	Windows	v. 2017.2	generate/write		generate/write	
Mnova	Mnova script exporting .mnova files	Damien Jeannerat	Any system running Mnova	V.1.1 (see below)	generate		generate (When spectra are available)	NMR records

Julien Wist
Niels Schlörer
Stefan Kuhn
Luc Patiny

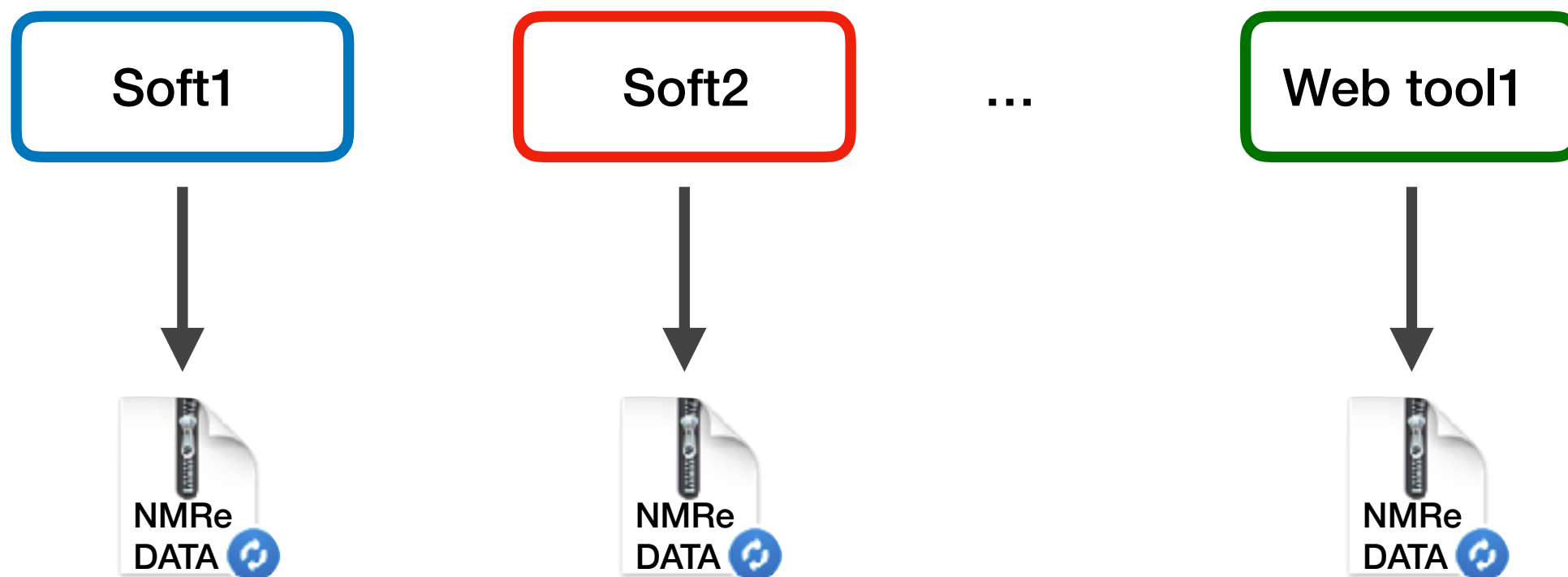
Database + visualisation + assignment tool + ...

Compatible software and web service

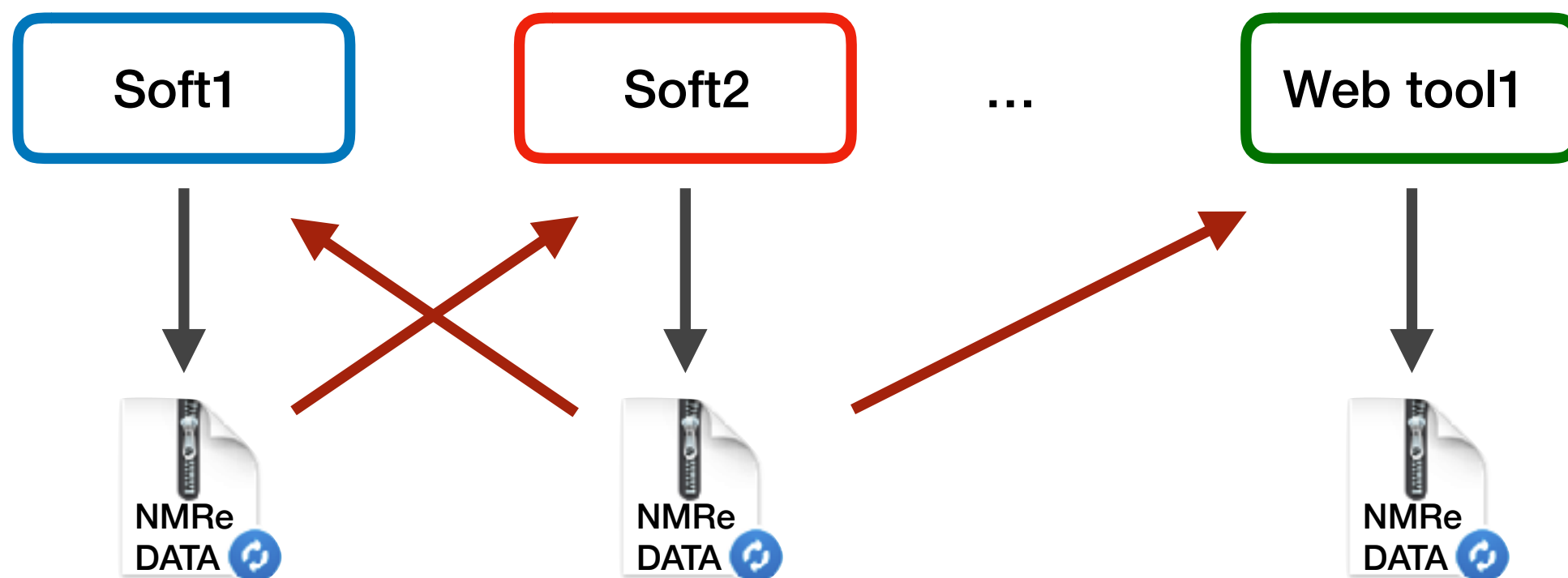
Announced Software & tools

Software Platform	Product/tool	Developer	Operating system	Version	NMReDATA	NMR Record	Examples of output
TopSpin	CMCse	Bruker	Windows, Linux and MAC OS X	V. ?	write	write	
Mnova	Mnova NMR	Mestrelab	Mac OS, Windows, Linux?	Announced for Q1 2018	?		
Matlab/Octave	NMReDATA_View	Damien Jeannerat	Any system running matlab/octave	V0 (will be on GitHub) when ready for test	read	read	
	NOMAD	Tomas Lebl					
Firefox/Google Chrome		Julien Wist	Any running Firefox/Google Chrome		generate	generate	

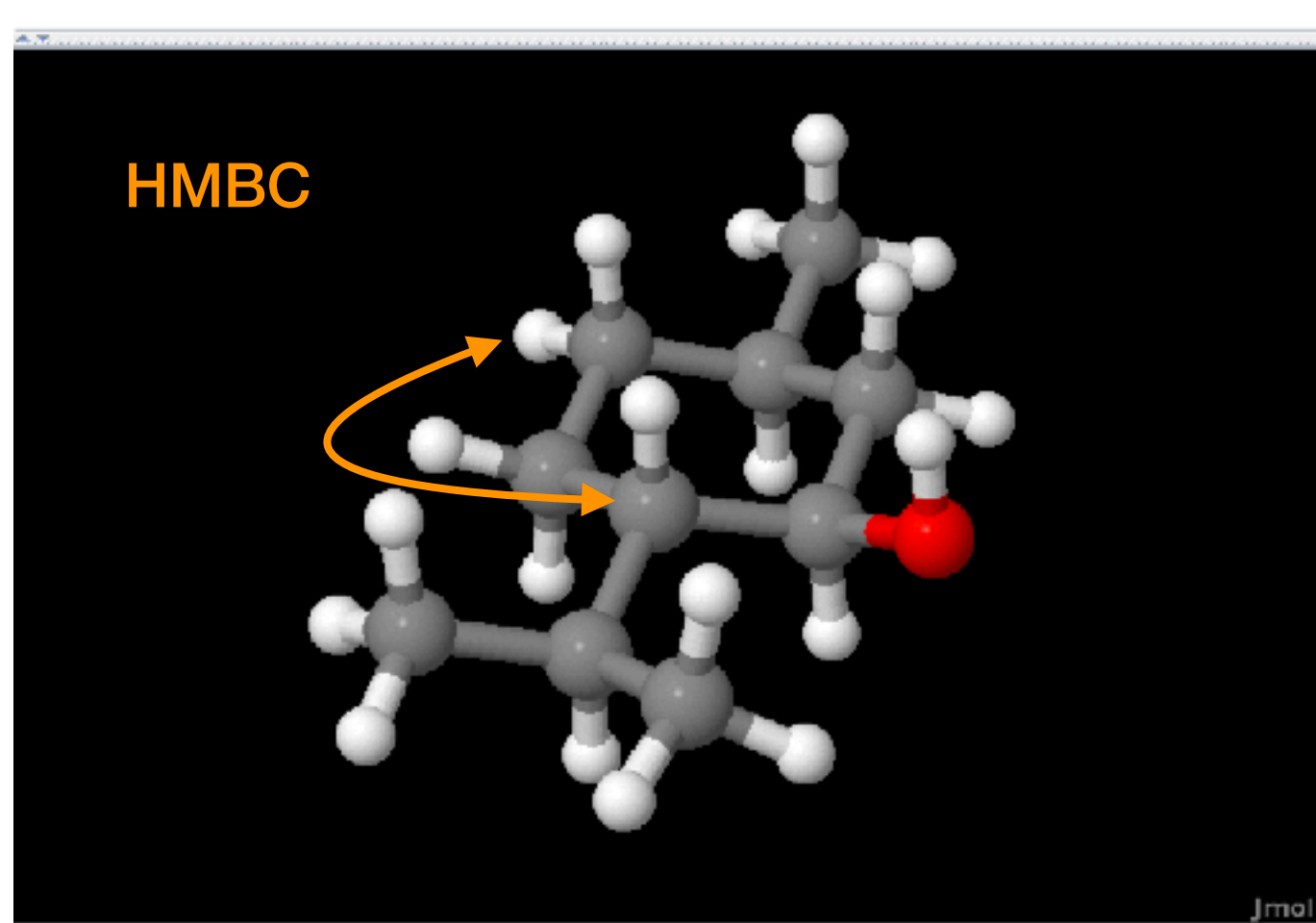
Need certification



Need certification



Need Visualizer/Validator

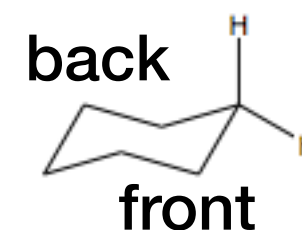


Need tutorial

Explicit H when possible/known



Watch for “conventional” explicit information

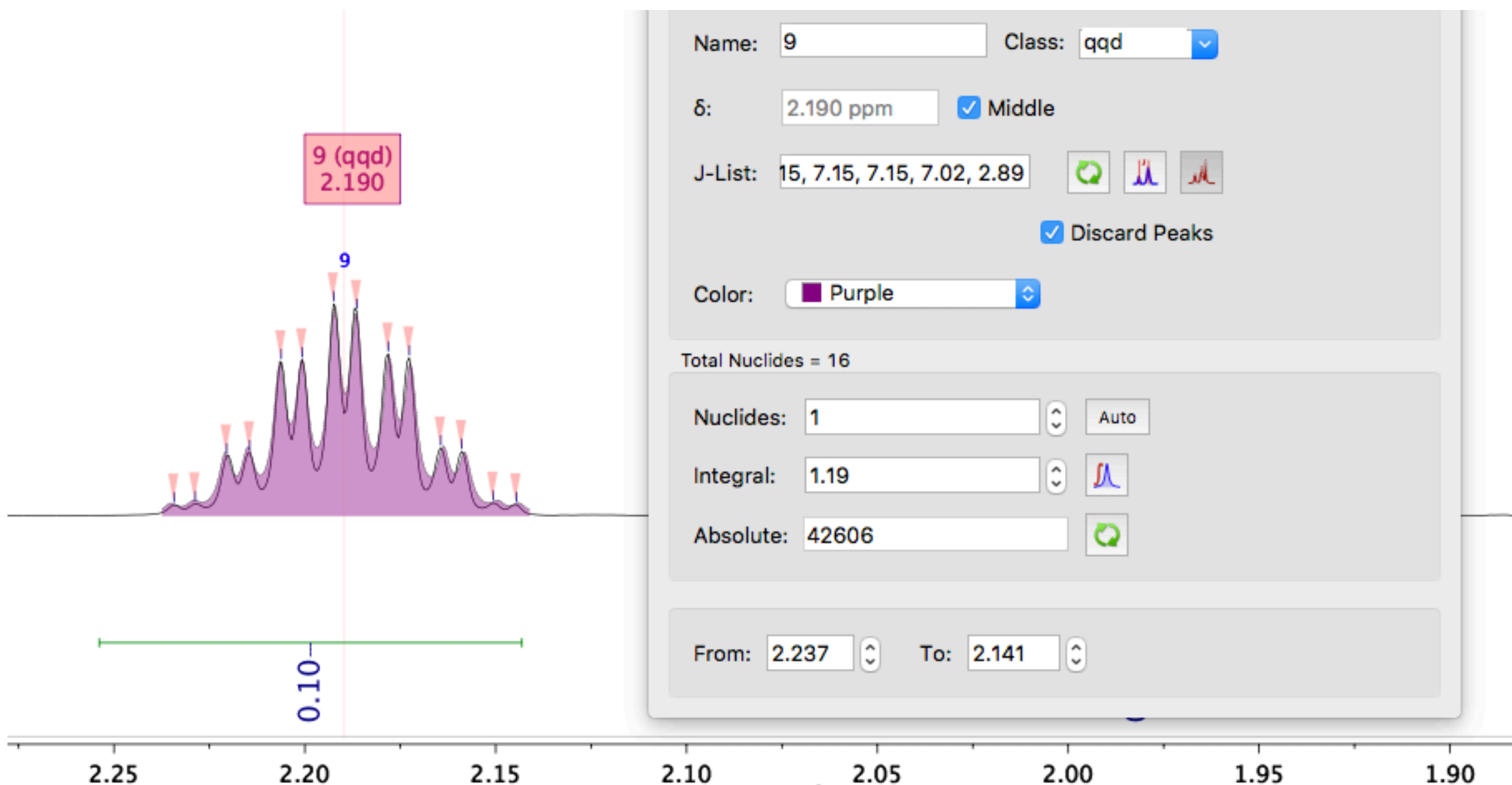


```

dj-air:3 djeanner$ cat ~/dropbox/demo.mol
demo.mol.sb-4f25a5d6-lDQsBy
ChemDraw07041811372D

  8  8  0  0  0  0  0  0  0  0  0999 V2000
    -0.9467   -0.0553    0.0000 C    0  0  0  0  0  0  0  0  0  0  0  0
    -1.3592   -0.7697    0.0000 C    0  0  0  0  0  0  0  0  0  0  0  0
    -0.5659   -0.5430    0.0000 C    0  0  0  0  0  0  0  0  0  0  0  0
     0.2321   -0.7697    0.0000 C    0  0  0  0  0  0  0  0  0  0  0  0
     0.6447   -0.0553    0.0000 C    0  0  0  0  0  0  0  0  0  0  0  0
    -0.1486   -0.2819    0.0000 C    0  0  0  0  0  0  0  0  0  0  0  0
     0.6447     0.7697    0.0000 H    0  0  0  0  0  0  0  0  0  0  0  0
     1.3592   -0.4678    0.0000 H    0  0  0  0  0  0  0  0  0  0  0  0
  1  2  1  0
  2  3  1  0
  3  4  1  0
  4  5  1  0
  5  6  1  0
  6  1  1  0
  5  7  1  0
  5  8  1  0
M  END
  
```

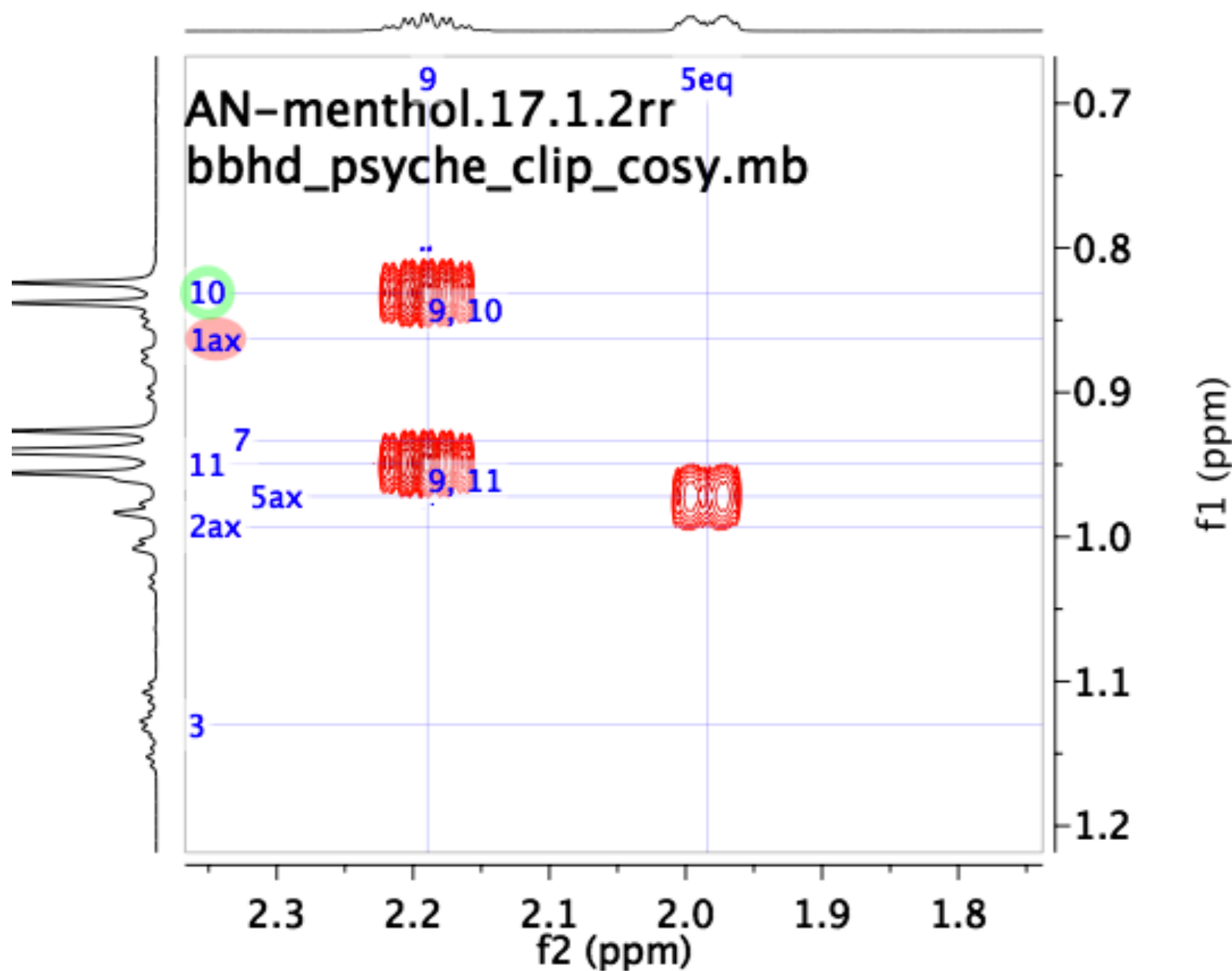

Improve work when exploiting spectra



Improve work when exploiting spectra

Atom	Chemical	Quality	COSY	Predicted	J	NOE	HSQC
1 C	34.57						1b, 1a, 1eq, 1ax
1a...	0.86	-0.31			12.75(...		1
1e...	1.68		☒ 1...		12.75(...		
2 C	23.14						2a, 2eq, 2ax
2a...	0.99		☒ 1...		3.30(1...		2
2e...	1.63		☒ 1...		3.20(1...		2
▼ 3 C	50.16						3
H	1.13		☒ 4		12.76(...		3
▼ 4 C	71.59						4
H	3.43		☒ 1...		9.90(...		4
5 C	45.06						5b, 5a, 5eq, 5ax
5a...	0.97		☒ 4		10.90(...		5
5e...	1.98		☒ 4		4.50(...		5
▼ 6 C	31.62						6
H	1.44		☒ 1...		6.58(...		6
▼ 7 C	22.23						7
	0.93				6.58(6)		7
▼ 9 C	25.84						9
H	2.19		☒ 1...		2.72(3...		9
▼ 10 ...	16.10						10
	0.83		☒ 9		6.98(9)		10
▼ 11 C	20.99						11
	0.95		☒ 9		7.05(9)		11
▼ 12 ...							
H	1.33..1...		☒ 4		4.80(4)		

Improve work when exploiting spectra



IUPAC, CODATA, FAIRness



IUPAC INTERNATIONAL UNION OF PURE AND APPLIED CHEMISTRY

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16 JULY 2018 - 17 JULY 2018

SUPPORTING FAIR EXCHANGE OF CHEMICAL DATA THROUGH STANDARDS DEVELOPMENT

This event has passed.

JCAMP DX/&/?

FAIR

Spectra :
 Manufacturer's format
 Jcamp (simulations, be FAIR)

DATE & TIME

VENUE
 University of Amsterdam, Science Park
 Science Park
 Amsterdam, Netherlands
 + Google Map

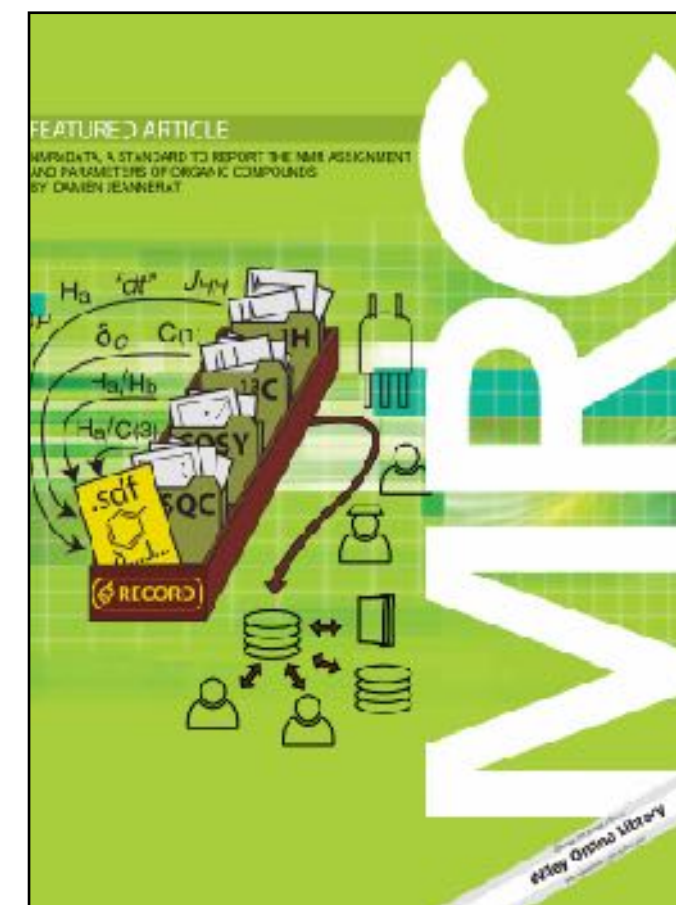
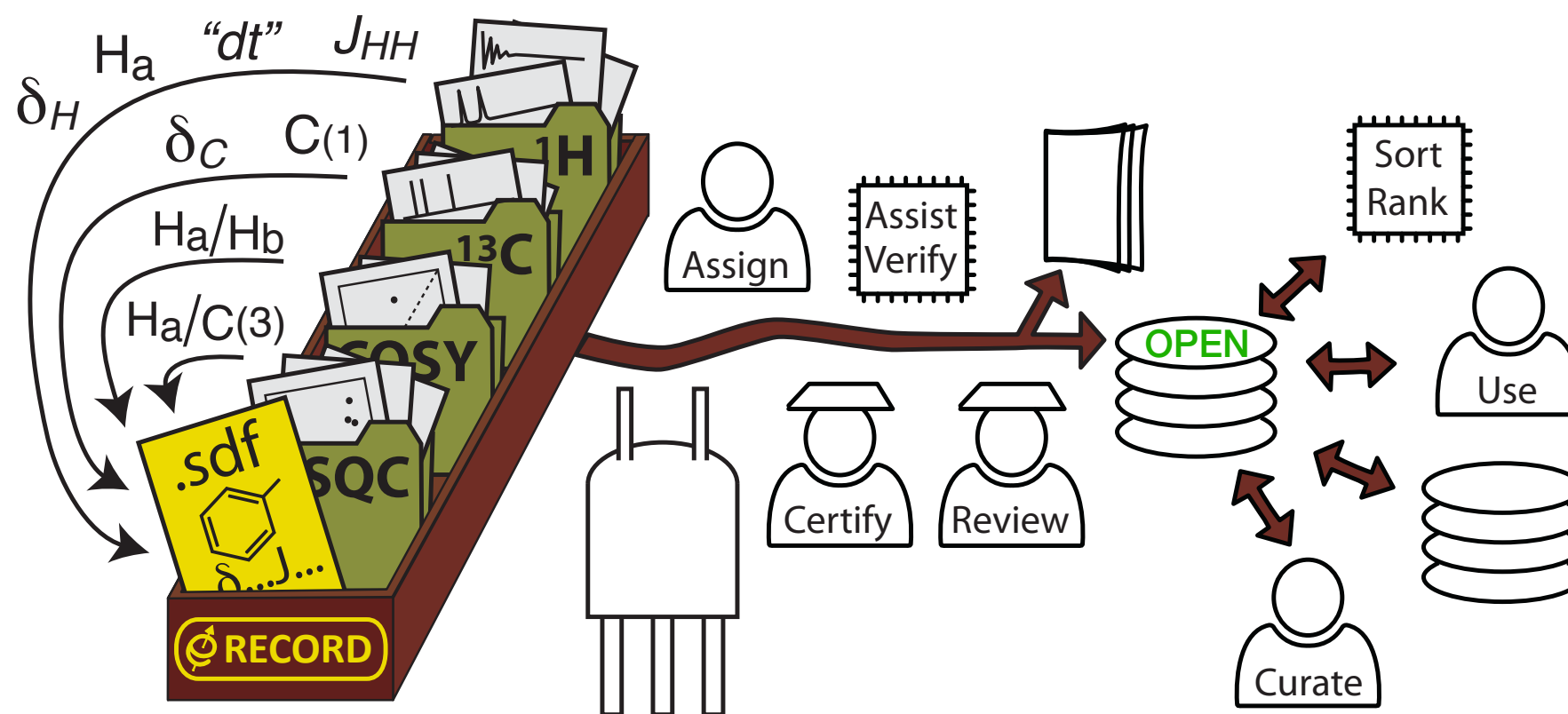
Twitter Facebook Google+ LinkedIn +

Conclusion

Momentum

Board of advisers / deciders

Support / Certification (free for non-profit....)



Pupier, M.; Nuzillard, J.-M.; Wist, J.; Schlörer, N. E.; Kuhn, S.; Erdelyi, M.; Steinbeck, C.; Williams, A. J.; Butts, C.; Claridge, T. D. W.; Mikhova, B.; Robien, W.; Dashti, H.; Eghbalian, H. R.; Farès, C.; Adam, C.; Kessler, P.; Moriaud, F.; Elyashberg, M.; Argyropoulos, D.; Pérez, M.; Giraudeau, P.; Gil, R. R.; Trevorow, P.; Jeannerat, D., NMReDATA, a standard to report the NMR assignment and parameters of organic compounds. *Magn. Reson. in Chem.* **2018**.

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Christoph Steinbeck⁶, Antony J. Williams^{9,&}, Craig Butts¹⁰, Bozhana Mikhova^{11,*}, Damien Jeannerat^{4,§}

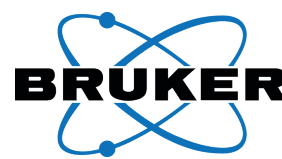
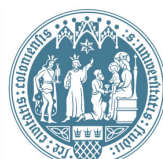
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