

# NMReDATA software and nmrshiftdb2

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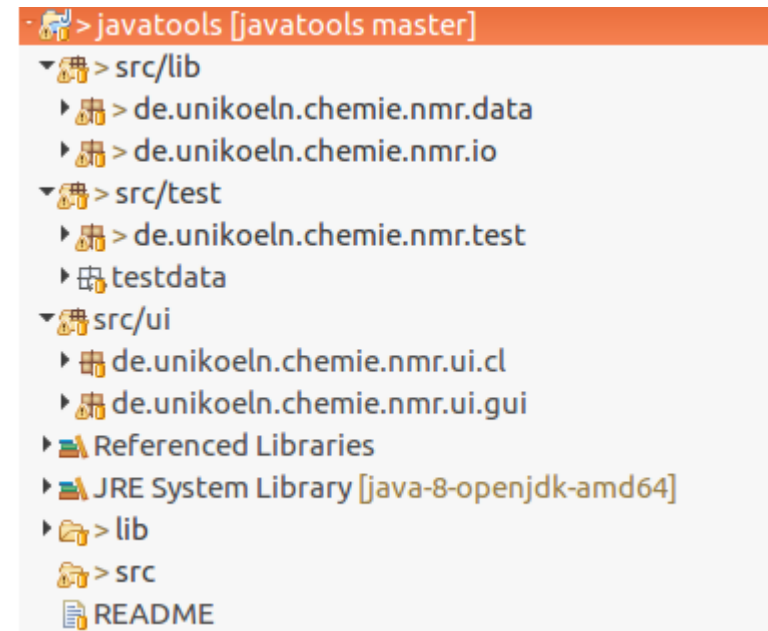
# Overview

- The NMReDATA tools
- Standalone functions
- Use as library
- Example: LDS
- nmrshiftdb2

# The Java tools library

- NMRReDATA java tools contain a java library for reading and writing NMRReDATA files
- Internal object come from the jcamp-dx library
- Allows direct reading/writing as jcamp
- Programs can integrate the library and navigate the object tree
- Also serves as conformity check

<https://github.com/NMRReDATAInitiative/javatools>



# The Java tools library

- Offers command line and GUI interface
- GUI allows opening, viewing and saving as NMReDATA, LSD or jcamp, editing and more data formats will follow

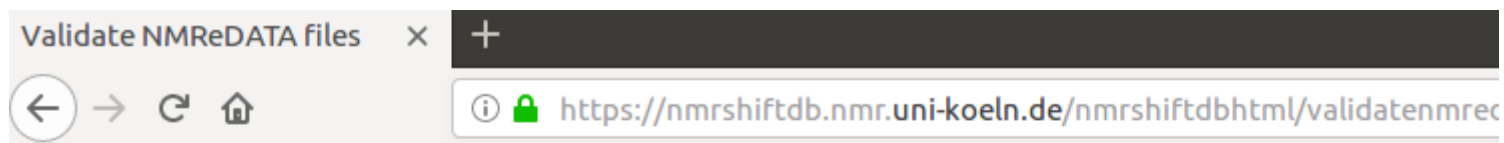
The screenshot displays the NMReData editor interface. The main window shows a chemical structure of an alkene derivative with atoms labeled 1 through 8. The structure is CC=CC(=O)OCC. The labels are: 1 (C=O), 2 (alkene C), 3 (alkene H), 4 (ethyl CH2), 5 (ethyl CH3), 6 (ethyl CH3), 7 (O), and 8 (ether O). The table on the right lists the following data:

Shift	Intensity	Assignn
1.255	0.0	6
1.854	0.0	5
4.177	0.0	4
5.844	0.0	3
6.949	0.0	2

Below the table, there are input fields for 'Frequency' (set to 500) and 'Spectrum\_Location' (set to file:).

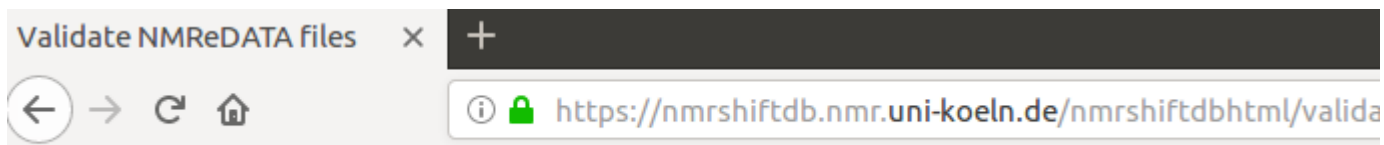
# The Java tools library

- Integration into web applications is possible
- Online check is at <https://nmrshiftdb.nmr.uni-koeln.de/nmrshiftdbhtml/validatenmredata.jsp>



Select NMReDATA file to upload:  No file selected.

Uploaded Filename: cmcse.sdf  
The molecule in your file has formula C<sub>6</sub>H<sub>10</sub>O<sub>2</sub>  
Your file contains 5 spectra  
Spectrum 1 has 5 peaks  
Spectrum 2 has 6 peaks



Select NMReDATA file to upload:  No file selected.

Uploaded Filename: example.nmredata.sdf  
version and level are compulsory!

# The Java tools library

- Can be integrated in other products
- Example: LSD
- Has its own format, not very complicated
- 65 lines of code in Java tools make them write LSD files

```
C:\Users\Privat#\Desktop\LSDos-3.4.10>lsgnmredata.bat "HAP_benzo(a)pyrene_assignments_1.nmredata.sdf"
```

```
Converting NMReDATA file to LSD file...
```

```
...
```

```
1 solution found.
```

```
Result written to "HAP_benzo(a)pyrene_assignments_1.nmredata.sdf".sol.sdf
```

```
C:\Users\Privat#\Desktop\LSDos-3.4.10>lsgnmredata.bat cmcse.sdf
```

```
Converting NMReDATA file to LSD file...
```

```
...
```

```
1 solution found.
```

```
Result written to cmcse.sdf.sol.sdf
```

# nmrshiftdb2

NMReDATA (by using the Java tools) has been integrated into nmrshiftdb2 for the following functions:

- Loading data into QuickCheck and assignment tool
- Submitting data
- Downloading data (original file as well as generated, NMR Record is possible as well)
- Downloading the database as NMReDATA

# Nmrshiftdb2 - QuickCheck

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:  cmcse.sdf

Atom	$\delta$ [ppm]	Deviation from prediction
1	166.603	0.30
2	144.493	0.34
3	122.699	0.35
4	60.151	0.25
5	17.98	0.08
6	14.166	0.07

Home Search Results **Quick Check** Predict Assignment Submit Review Help

Quick Check  
Note that this feature is currently experimental!

...

1D spectra 2D spectra

Atom No. ↓ <sup>13</sup>C Shift

1     
 ●●● 166.30, diff: 0.30

2     
 ●●● 144.15, diff: 0.34

3     
 ●●● 123.05, diff: 0.35

4     
 ●●● 59.90, diff: 0.25

5     
 ●●● 17.90, diff: 0.08

6     
 ●●● 14.24, diff: 0.07

<sup>1</sup>H Shift

H9     
 ●●● 6.95, diff: 0.00

H10     
 ●●● 5.82, diff: 0.02

H11     
 ●●● 4.16, diff: 0.02

H13     
 ●●● 1.86, diff: 0.01

H16     
 ●●● 1.22, diff: 0.04

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

[Submit <sup>13</sup>C](#)

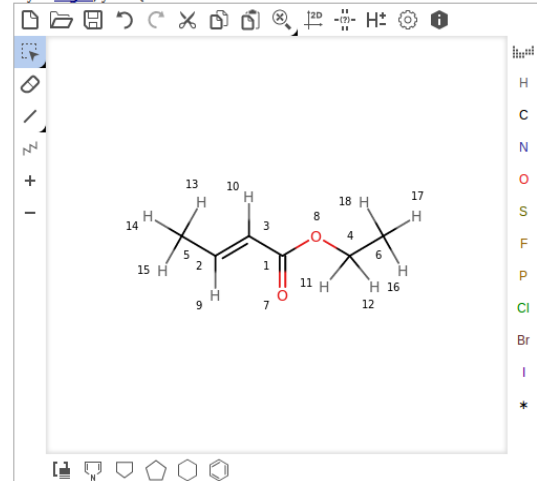
Input list: [Input format](#)

[Submit <sup>1</sup>H](#)

Input list: [Input format](#)

Use .SHIFT REFERENCE in jcamp files (e. g. files from Topspin)

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

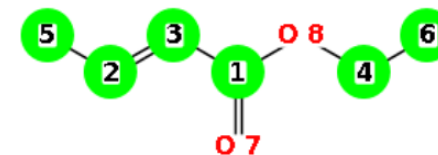


Choose an MNova mol or sd file (MNova 11 or newer) or ChemDraw mol file with labels:

No file selected.

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

Choose an NMReDATA file:  No file selected.



10: nmrshiftdb2 quality check: accept/reliability: excellent



# Nmrshiftdb2 – Submitting data

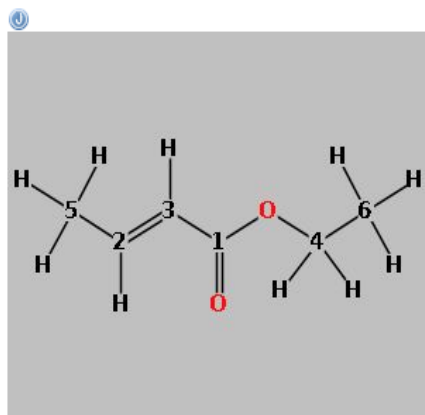
Choose an MNova mol or sd file (MNova 11 or newer) or ChemDraw mol file with labels:  No file selected.

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

Choose an NMRReDATA file:  cm



Choose your spectrum type!



Spectrum type: 13C

Atom No. ↓	Shift	Mult. (coupling const.)
1	166.603	
2	144.493	
3	122.699	
4	60.151	
5	17.98	
6	14.166	

Chemical name(s):  
Double Bond Specification:  
CAS number:  
Conditions:  
Additional comments:  
Additional spectrum information:  
Spectrum keywords:  
No JCAMP-DX file attached!  
No PDF file attached!  
No raw data file attached!  
No image file attached!  
No other files attached!

Export your submit (does not save or finish the submit):

Get this molecule as  file

Get this spectrum as  file

Get this spectrum and its molecule as

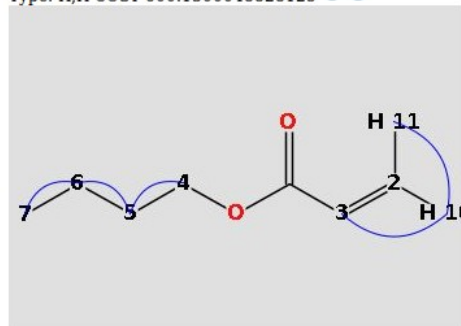
Your Mnova or Topspin file contains the spectrum types given below. Choose which one you want to submit now. You submit the other spectrum by uploading the file again, if you have not yet done both! Spectrum type:

# Nmrshiftdb2 – Downloading data

General chemie koeln RUB/CDC13

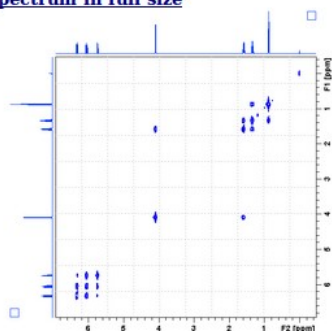
Spectral Data Additional Data Download

Type: H,H-COSY 600.1300048828125



Axis: 1 (H)		Axis: 2 (H)	
Atom	Shift	Atom	Shift
6	1.34	7	0.88
5	1.59	6	1.34
4	4.09	5	1.59
2	6.32	3	6.05
2	6.32	2	5.74

Show spectrum in full size



...

[Report incorrect data](#)

Back

[Copy dataset link](#)











Get this dataset as  file

[Copy molecule link](#)

Submit a new  molecule

- NMR record
- NMReDATA
- cml
- Report

# Nmrshiftdb2 – Downloading the database (available soon)

File	Date	Author	Commit
<a href="#">.project</a>	2015-09-05	 shk3	<a href="#">[r948]</a> new predictor jars
<a href="#">nmrshiftdb2.sd</a>	2019-04-25	 shk3	<a href="#">[r1876]</a> 2019-04-25
<a href="#">nmrshiftdb2.xml</a>	2019-04-25	 shk3	<a href="#">[r1876]</a> 2019-04-25
<a href="#">nmrshiftdb2_3d.sd</a>	2019-04-25	 shk3	<a href="#">[r1876]</a> 2019-04-25
<a href="#">nmrshiftdb2_3d.xml</a>	2019-04-25	 shk3	<a href="#">[r1876]</a> 2019-04-25
<a href="#">nmrshiftdb2datalicense.txt</a>	2015-09-01	 shk3	<a href="#">[r942]</a> 2015-09-01
<a href="#">nmrshiftdb2withsignals.sd</a>	2019-04-25	 shk3	<a href="#">[r1876]</a> 2019-04-25
<a href="#">nmrshiftdb2withsignals_3d.sd</a>	2019-04-25	 shk3	<a href="#">[r1876]</a> 2019-04-25
<a href="#">predictorc.jar</a>	2019-07-30	 shk3	<a href="#">[r1929]</a> predictor jars with 3d hose codes
<a href="#">predictorh.jar</a>	2019-07-30	 shk3	<a href="#">[r1929]</a> predictor jars with 3d hose codes

# Conclusions

- The NMReData Java tools offer a possibility to view, check, and convert NMReDATA files
- They can be integrated easily into other programs
- By integrating them into nmrshiftdb2, NMReDATA can be imported and exported to nmrshiftdb2
- This enables easy integration of data sources