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Title: Programs for 2D J prediction

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URL: <http://dmar.riken.jp/NMRinformatics/>

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Reference: Ito, K., Xu, X., and Kikuchi, J. "Improved prediction of carbonless NMR spectra by the machine learning of theoretical and fragment descriptors for environmental mixture analysis" (submitted)

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About

Programs for prediction of 2D J -resolved NMR spectroscopy by combination approach with quantum mechanical computation and machine learning, including the machine learning input file generator, predicted 2D J -resolved NMR spectra generator, and predicted J -coupling tree drawer, and so on.

Requirements

1. [Python](#)
[rdkit](#)
[openbabel](#)
[nmrglue](#)
2. [R](#)
[caret](#)
3. [Java](#)
4. [Node.js](#)
[openchemlib-extended](#)
5. [Gaussian](#)

Example of using

(1) Quantum mechanical computation

[Data and programs: 1_QM directory in [Predict 2DJ programs and data.zip](#)]

1-1. Execute command for DFT calculation

```
g16 33032.com
```

1-2. Convert Gaussian log files to sdf and mol files

```
cd your\_file\_path/1_QM/
```

```
python ob_conv.py your_file_path/1_QM
```

(2) The machine learning input file generation of $^nJ_{\text{HH}}$ coupling constant for training dataset

[Data and programs: 2_JtrainDATA directory in [Predict 2DJ programs and data.zip](#)]

2-1. Execute python program

```
cd your_file_path/2_JtrainDATA/  
python Jtrain.py
```

2-2. GUI input arguments

- ① your_file_path/2_JtrainDATA/
- ② input_Jtrain.xlsx
- ③ your_file_path/1_QM/TrainData/log
- ④ your_file_path/1_QM/TrainData/sdf
- ⑤ your_file_path/1_QM/TrainData/mol
- ⑥ your_openchemlib-extended_path/node_modules

(3) The machine learning input file generation of ^1H chemical shift for training dataset

[Data and programs: 3_CStrainDATA directory in [Predict 2DJ programs and data.zip](#)]

3-1. Execute python program

```
cd your_file_path/3_CStrainDATA/  
python CStrain.py
```

3-2. GUI input arguments

- ① your_file_path/3_CStrainDATA/
- ② input_CStrain.xlsx
- ③ your_file_path/1_QM/TrainData/log
- ④ your_file_path/1_QM/TrainData/sdf
- ⑤ your_file_path/1_QM/TrainData/mol
- ⑥ your_openchemlib-extended_path/node_modules

(4) The machine learning input file generation of $^nJ_{\text{HH}}$ coupling constant for test dataset

[Data and programs: 4_JtestDATA directory in [Predict 2DJ programs and data.zip](#)]

4-1. Execute python program

```
cd your_file_path/4_JtestDATA/  
python Jtest.py
```

4-2. GUI input arguments

- ① your_file_path/4_JtestDATA/
- ② input_Jtest.xlsx
- ③ your_file_path/1_QM/TestData/log
- ④ your_file_path/1_QM/TestData/sdf
- ⑤ your_file_path/1_QM/TestData/mol
- ⑥ your_openchemlib-extended_path/node_modules

(5) The machine learning input file generation of ^1H chemical shift for test dataset

[Data and programs: 5_CStestDATA directory in [Predict 2DJ programs and data.zip](#)]

5-1. Execute python program

```
cd your_file_path/5_CStestDATA/  
python CStest.py
```

5-2. GUI input arguments

- ① your_file_path/5_CStestDATA/
- ② input_CStest.xlsx
- ③ your_file_path/1_QM/TestData/log
- ④ your_file_path/1_QM/TestData/sdf
- ⑤ your_file_path/1_QM/TestData/mol
- ⑥ your_openchemlib-extended_path/node_modules

(2~5) Common GUI setting

J-coupling Prediction Input Helper

Information needed to produce the processed input file

Directory path of the basic input ①

Name of the basic input file: ②

Directory path of the LOG files ③

Directory path of the SDF files: ④

Directory path of the MOL files ⑤

Path of node modules(for OpenChemLib) ⑥

Produce

(6) Generate fragment descriptors

[Data and programs: 6_GenFra directory in [Predict 2DJ programs and data.zip](#)]

6-1. Execute batch program

```
cd your_file_path/6_GenFra/CS_train/water/  
./toolgaussianlearndata.bat  
cd your_file_path/6_GenFra/CS_train/methanol/  
./toolgaussianlearndata.bat  
cd your_file_path/6_GenFra/CS_test/water/  
./toolgaussianlearndata.bat
```

More details are shown in [previous study](#).

(7) Machine learning of ${}^nJ_{\text{HH}}$ coupling constant

[Data and programs: 7_Jlearning directory in [Predict 2DJ programs and data.zip](#)]

7-1. Execute R program

```
cd your\_file\_path/7_Jlearning/
```

```
Rscript ML_Pred_tool_CMD.R Core=2 K_CV=5 Grid=3  
ML_Numbers=36,48,145,169,173,191-192,236-237 outFile_Name=J_Train  
inFile_Path=your\_file\_path/7_Jlearning/ML_input_J_train.csv
```

(8) Machine learning of ^1H chemical shift

[Data and programs: 8_CSlearning directory in [Predict 2DJ programs and data.zip](#)]

8-1. Execute R program

```
cd your\_file\_path/8_CSlearning/
```

```
Rscript ML_Pred_tool_CMD.R Core=2 K_CV=5 Grid=3  
ML_Numbers=36,48,145,169,173,191-192,236-237 outFile_Name=CS_Train  
inFile_Path=your\_file\_path/8_CSlearning/ML_input_CS_train.csv
```

(9) Prediction of $^nJ_{\text{HH}}$ coupling constant

[Data and programs: 9_Jpred directory in [Predict 2DJ programs and data.zip](#)]

9-1. Execute R program

```
cd your\_file\_path/9_Jpred/
```

```
Rscript ML_Test_tool_CMD.R Rdata_Path=your\_file\_path/7_Jlearning/J_Train.Rdata  
test_File_Path=your\_file\_path/9_Jpred/ML_input_J_test.csv test_Out_Name=J_Test
```

(10) Prediction of ^1H chemical shift

[Data and programs: 10_CSpred directory in [Predict 2DJ programs and data.zip](#)]

10-1. Execute R program

```
cd your\_file\_path/10_CSpred/
```

```
Rscript ML_Test_tool_CMD.R Rdata_Path=your\_file\_path/8_CSlearning/CS_Train.Rdata  
test_File_Path=your\_file\_path/10_CSpred/ML_input_CS_test.csv test_Out_Name=CS_Test
```

(11) Predicted 2D J -resolved NMR spectra generation

[Data and programs: 11_2DJgen directory in [Predict 2DJ programs and data.zip](#)]

11-1. Execute python program

```
cd your\_file\_path/11_2DJgen/
```

```
python plot2DJ.py CSpred.csv Jpred.csv your\_file\_path/11_2DJgen/999/pdata/
```

(12) Draw the predicted J coupling tree

[Data and programs: 12_JtreeDraw directory in [Predict 2DJ programs and data.zip](#)]

12-1. Execute python program

```
cd your\_file\_path/12_JtreeDraw/
```

```
python plotTree.py your\_file\_path/11_2DJgen/TreeTable.csv 1
```