week2b

August 23, 2023

Notebook file for Week 2, T2 Relaxation experiment

Import libraries. The nmrbase folder needs to be located within the folder containing this Jupyter notebook

```
[]: import numpy as np
import scipy
import matplotlib.pyplot as plt
import nmrbase.expbase as expbase
import nmrbase.expdta as expdta
```

T2 Relaxation NOTE: To analyze and report multiple datasets, simply copy the necessary blocks of code.

```
[]: filename = r''../DIRECTORY/FILENAME" #Defines the path to the data file. Here \rightarrow the path is relative to the current folder.
```

```
[]: a = expbase.expbase()
a.load(filename) # load data

f1=plt.figure()
ax1=f1.subplots()
a.split() # splitting raw T2 data for later
→processes
a.plottm(ax1,0)
```

```
[]: ## TASKS:
    ## fine tune with set_xlim and set_ylim parameters to zoom in on the echo
    ax1.set_ylim([VALUE, VALUE])
    ax1.set_xlim([VALUE, VALUE])
```

```
[]: a.pproc['digfmin']=1500  # set appropriate digital filter parameters
a.pproc['digfmax']=3500
a.digfilt()  # perform digital filter operation
```

```
[ ]: ## TASKS:
     ## fine tune with set_xlim and set_ylim parameters to zoom in on the echo
     f2=plt.figure()
     ax2=f2.subplots()
     a.plottm(ax2,0) # plot the entire scan
     ax2.set_ylim([VALUE, VALUE])
     ax2.set_xlim([VALUE, VALUE]) #zoom in on the 1st echo, which is the strongest one
[ ]: ## TASKS:
     ## change parameter "ftmin" and "ftmax" to select the time interval
     ⇔corresponding to the echoes for the Fourier transform
     ## change parameter "ffmin" and "ffmax" to select the frequency range of the NMR_1
      \rightarrow signal
     ## include statements to label axes
     ## use statements to change appearance such as font size, etc.
     f3=plt.figure()
     ax3 = f3.subplots()
     a.pproc['ftmin']=VALUE
                                                # time interval for Fourier transform
     \hookrightarrow (0 = all data)
     a.pproc['ftmax']=VALUE
     a.pproc['ffmin']=VALUE
                                               # frequency interval for spectrum_
     \rightarrow display (0 = all data)
     a.pproc['ffmax']=VALUE
     a.pproc['dispper']=0.5
                                           # leave 50% space between each acquired_
      \hookrightarrow spectrum
     a.proc()
                                             # calculate Fourier transform of the data
     ## this will use the digitally filtered data from before. Instead, the original \Box
      → data can be processed by loading it again.
     a.plotfrq(ax3,0)
                                            # plot the frequency domain data of the
      \rightarrow first scan from the data set
[]: # Time-resolved integration of peaks on NMR spectra
     ## TASKS:
     ## change the intmin and intmax parameters to select the correct frequency range,
      \rightarrow for integration
     ## use set_xlabel and set_ylabel to set the labels to get a publication quality.
      \hookrightarrow figure
```

```
f4=plt.figure()
ax4=f4.subplots()
                          # SET correct frequency range for integration
a.pproc['intmin']=VALUE
a.pproc['intmax']=VALUE
a.integrate()
                          # perform integration
#find the waiting time tau, then set the correct x-axis
dx=a.pproc["soff"]
x0=a.pproc["slen"]
print('x0 = ',x0,'s , dx = ',dx,'s')
                                          # x0 is the time of 1st echoes, and
\rightarrow dx is the interval between echoes
a.idt.x0=x0
a.idt.dx=dx
a.idt.plot(ax4,disp=[0])
                                         # disp=0 plots only the NMR signal in_{ij}
\rightarrow trace 0
ax4.set_xlabel("seconds [s]")
ax4.set_ylabel("Y LABEL")
```

Fitting of the integral to time axis. NOTE: For some results to fit, optimize the initial conditions

```
[]: # Fitting of the integral
     ## TASKS:
     ## change the initial fitting parameters to obtain proper fitting
     ## report the fitted relaxation constants with proper significant figures
     ## use set_xlabel and set_ylabel to set the labels to get a publication quality_
     \hookrightarrow figure
     ## adjust appearance of figure as needed
     ax5 = plt.figure().subplots()
     a.idt.plot(ax5,disp=[0])
                                                 # to be overlayed by fitted curve,
     \rightarrow disp=0 plots only the NMR signal in trace 0
     def fun(t,a,b,c):
         return FORMULA
                                                 # INPUT the formula used for fitting.
      \rightarrow Use "np.exp()" for exponential, and t for time-axis.
     y = a.idt.dta[0] # Integral data points to fit
     x = np.linspace(a.idt.x0,a.idt.x0+a.idt.dx*round(len(y)-1),len(y)) # Time axis
     a.p1,a.p2=scipy.optimize.
      →curve_fit(fun,x,y,p0=[A_VALUE,B_VALUE,C_VALUE],maxfev=5000)
                                                                                    #__
      \hookrightarrow Select the initial conditions
```