week3

June 15, 2023

Notebook file for Week 3, Diffusion 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
```

Diffusion 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⊔

→ the path is relative to the current folder.
```

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

f1=plt.figure()
ax1=f1.subplots()
a.plottm(ax1,1)
```

```
[]: ## TASKS:
## fine tune with set_xlim and set_ylim parameters to zoom in on the echo
ax1.set_ylim([-0.15,0.15])
ax1.set_xlim([0.25,0.35])
f1
```

```
[]: 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(ax2,1) # plot the 1st scan
ax2.set_ylim([-0.15,0.15])
ax2.set_xlim([0.25,0.35])
```

```
[]: f3=plt.figure()
     ax3 = f3.subplots()
     a.pproc['ftmin']=0
                                            # time interval for Fourier transform (0 = ___
      \hookrightarrow all data)
     a.pproc['ftmax']=0
     a.pproc['ffmin']=0
                                             # frequency interval for spectrum display...
      \hookrightarrow (0 = all data)
     a.pproc['ffmax']=0
     a.proc()
                                             # calculate Fourier transform of the data
      \hookrightarrow in a
     ## 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_
      ⇔first scan from the data set
     ## TASKS:
     ## change parameter "ftmin" and "ftmax" to select the time interval_{\sqcup}
      →corresponding to the echoes for the Fourier transform
     ## change parameter "ffmin" and "ffmax" to select the frequency range of the
      →NMR signal
     ## include statements to label axes
     ## use statements to change appearance such as font size, etc.
```

```
[]: | # Gradient-duration-resolved integration of peaks on NMR spectra
     f4=plt.figure()
     ax4=f4.subplots()
     a.pproc['intmin']=0
                             # set correct frequency range for integration
     a.pproc['intmax']=0
     a.integrate()
                                # perform integration
     #find the starting gradient duration and increment, then set the correct x-axis
     dx=a.pinc["inc"][0]
     x0=a.p["p2"]
     print('x0 = ',x0,'s, dx = ',dx,'s')
                                                # x0 is the starting gradient
      \hookrightarrow duration, and dx is the increment
     a.idt.x0=x0
     a.idt.dx=dx
```

```
a.idt.plot(ax4,disp=[0])  # disp=0 plots only the NMR signal_
in trace 0

ax4.set_xlabel("seconds [s]")
ax4.set_ylabel("Y LABEL")

## TASKS:
## change the intmin and intmax parameters to select the correct frequency_
range for integration

## use set_xlabel and set_ylabel to set the labels to get a publication quality_
rigure
```

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

```
[]: # Fitting of the integral
     ax5 = plt.figure().subplots()
     a.idt.plot(ax5,disp=[0])
                                               # to be overlayed by fitted curve,
     →disp=0 plots only the NMR signal in trace 0
     big_delta=a.p.get('tau',0.1)
                                                 # time interval between gradients
     def fun(t,a,b):
        return FORMULA
                                                 # INPUT the formula used for
      ofitting. Use "np.exp()" for exponential, and t for small-delta (gradientum)
     \rightarrow duration).
     y = a.idt.dta[0]
     x = np.linspace(a.idt.x0,a.idt.x0+a.idt.dx*(a.pinc['n'][0]-1),a.pinc['n'][0])
     a.p1,a.p2=scipy.optimize.curve_fit(fun,x,y,p0=[y[0],0.001],maxfev=5000)
                # OPTIMIZE the initial conditions, especially the exponential index
     print(round(a.p1[1],SF1), '±', round(np.linalg.eig(a.p2)[0][1]**0.5,SF2))
                   # REPLACE "SF1" and "SF2" with positive integrers to report the \Box
     ⇔fitting results with correct significant figures
     x=np.linspace(x[0],x[-1],1000)
                                                              # use 1000 points to
     ⇒generate a smooth curve
     a.fit_points = a.p1[0]*np.exp(-a.p1[1]*x**2*(big_delta-x/3))
     pl,=ax5.plot(x,a.fit_points,'r-')
     ax5.set_xlabel("seconds [s]")
                                                         # SET the labels to get a_{\square}
     →publication quality figure
     ax5.set ylabel("Y LABEL")
     pl.figure.set_tight_layout('pad')
     pl.figure.canvas.draw()
```

```
## TASKS:

## change the initial fitting parameters to obtain proper fitting

## report the fitted gradient strength/diffusion coefficient with proper_

significant figures

## use set_xlabel and set_ylabel to set the labels to get a publication quality_

figure

## adjust appearance of figure as needed
```