## Script for the Correction of differential scattering cross section in SAXS experiments (cooking book)

```
In [1]: import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
```

URL for SAXS data of glassy carbon reference. There are several different onex available. The signal in high Q is nearly the same for all of them. Only in the low Q region they diverge from one another. The urls shown below correspond to the Glassy\_Carbon\_A, \_B and \_C samples from ANL. This script uses a Glassy carbon sample measured in another institute as reference.

Function to correct the thickness of the sample, by measuring an empty cuvette and one filled with water. For that two measurements with transmission and empty beam correspondingly have to be done with H2O and an empty cuvette. The mass attenuation coefficient for water in room temperature is

 $rac{\mu}{
ho}=10.37rac{cm^{z}}{g}$  . The sample thickness d can then be calculated by using: \vspace{3mm}

$$d = -
ho_{H_2O} rac{
ho}{\mu} ln \left(rac{T_{H_2O+cell}}{T_{cell}}
ight)$$

\vspace{3mm} , with T being the transmission factors for an empty cell and water and  $ho_{H_2O}=1rac{g}{cm^2}.$ 

%display latex Function to calculate the transmission factor for intensity of sample S and empty beam intensity  $I_0$ : \vspace{3mm}

$$T_S = rac{I_S}{I_0}$$

```
In [3]: def calc_T(I_S,I_0):
    return(I_S/I_0)
```

This Function calulates the Scaling or Correction Factor with a Glassy Carbon measurement done in your setup and scales it so it fits the intensity of the 'Glassy-Carbon-A' sample from ANL. You can chose whatever reference sample you like. They should turn out more or less the same. For consistency's sake, I will solely use the A sample. The formula for the SF calculation follows from Fan-Paper: \vspace{3mm}

$$SF = rac{\left(rac{\delta \Sigma}{\delta \Omega}
ight)_{ref}}{rac{(I_{GC})}{d_{GC}t_{GC}T_{GC}}}$$

\vspace{3mm}, with 'st' corresponding to the standard sample, in my case glassy carbon. BG corresponds to the background measurement and T the transmission factor for the GC measurements.

```
In [13]:
         # Übergabe:
         #
              - Intensity of GC
         #
              - Q values of measurement for interpolation from Reference samp
              - Background of GC measurement
              - Transmissionfactor GC
              - thickness sample d GC
         def calc_SF(I_GC,Q_GC,T_GC,d_GC,t_GC,plot=False):
             from scipy.optimize import curve fit as fit
             def fit_lin(x,a,c):
                 return np.exp(a*np.log10(x)+c)
             import os
             #correct this line for the path in which the GC reference is loc
         ated
             sample GCA = np.loadtxt(r'C:\Users\ZechT\FAUbox\Auswertung\Softw
         are\ICSPy\compute\SAXS\reference/GC_APS.txt')
             # in this line the linear fits at low Q values are made for the
         GC measurement and reference
             DSC GC = fit(fit lin,Q GC[:6],I GC[:6]/(d GC*t GC*T GC))[0]
             DSC Ref = fit(fit lin, sample GCA[:35,0], sample GCA[:35,1])[0]
             print('Fit Params. Meas: ',np.exp(DSC_GC), ' Fit Params. Ref: ',
         np.exp(DSC Ref))
             # the scaling factor is calculated
             SF = np.exp(DSC Ref[1])/(np.exp(DSC GC[1]))
             print(SF)
             SF = np.mean(SF)
             if plot:
                 import matplotlib.pyplot as plt
                 ax = plt.figure().add subplot(111)
                 ax.set xscale('log')
                 ax.set yscale('log')
                 ax.plot(Q GC, fit lin(Q GC, DSC GC[0], DSC GC[1]), '--b')
                 ax.plot(Q GC,fit lin(Q GC,DSC Ref[0],DSC Ref[1]),'--k')
                 ax.plot(Q GC,I GC/(d GC*t GC*T GC),'-r',label='meas. GC')
                 ax.plot(Q GC,SF*I GC/(d GC*t GC*T GC),'-b',label='meas. GC')
                 ax.plot(sample GCA[:,0],sample GCA[:,1],'-k',label='meas. GC
          • )
             return (SF, DSC GC, DSC Ref)
```

Finally the differential scattering crosssection for sample s will be correctly calculated using all previously determined parameters SF and  $d_s$ : \vspace{3mm}

$$\left(rac{\delta \Sigma}{\delta \Omega}
ight)_s(q) = SF\left(rac{I_s(q)}{d_s t_s T_s} - rac{BG_s(q)}{d_{BG} t_{BG} T_{BG}}
ight)$$

## From here on follows an example

From here follows an example using sample TiS710 and data measured on our Anton Paar machine. First the intensities are extracted from .stat files. This process will not be discussed further, because it's simply parsing the files contained in the in 'dir' specified folder automatically. This step corresponds to point (1) in the cooking book. Calculate the transmission factors. (You have calculated the Transmission by hand via Fit2D. You can therefore ignore the following code section. Important is only the following table in which all the transmission values for the data that is relevant in the following correction steps.)

```
In [26]: import os
         dir = r'C:\Users\ZechT\FAUbox\Auswertung\Software\ICSPy\compute\SAX
         S\test data\\'
         1 EB Stats = []
         1 TR Stats = []
         table = pd.DataFrame(columns = ['TR', 'EB', 'I TR', 'I EB', 'T'])
         pairs = pd.DataFrame(columns = ['typ', 'samp'])
         for iFile in os.listdir(dir):
             if os.path.isfile(os.path.join(dir, iFile)):
                 if '.stat' in iFile:
                      enum = iFile.split('_')[0]
                      typ = iFile.split(' ')[2]
                      samp = iFile.split(' ')[1]
                      pairs.loc[str(enum)] = [str(typ),str(samp)]
         for iFile in os.listdir(dir):
             if os.path.isfile(os.path.join(dir, iFile)):
                  #print(os.path.join(dir,iFile))
                 if '.stat' in iFile:
                      for line in open(os.path.join(dir,iFile),'r'):
                          if 'Total intensity' in line:
                              inty = line.split()[3]
                              enum = iFile.split(' ')[0]
                              samp = pairs.loc[str(enum), 'samp']
                              typ = pairs.loc[str(enum),'typ']
                              if str(samp) not in table.index:
                                  table.loc[str(samp)] = [0,0,0,0,0]
                              if typ == 'TR':
                                  table.loc[str(samp),'TR'] = enum
                                  table.loc[str(samp),'I_TR'] = inty
                              elif typ == 'EB' or 'DB':
                                  table.loc[str(samp),'EB'] = enum
                                  table.loc[str(samp),'I EB'] = inty
         for iInd in table.index:
             table.loc[iInd,'T'] = calc_T(float(table.loc[iInd,'I_TR']),float
          (table.loc[iInd,'I EB']))
         table
```

## Out [26]:

	TR	EB	I_TR	I_EB	Т
GC	01724	01723	3.75920E+05	7.01408E+05	0.535951
SGNR	01727	01728	2.11857E+05	7.99840E+05	0.264874
H2O	01744	01743	1.45425E+05	5.66515E+05	0.256701
EC	01746	01747	3.65022E+05	5.57136E+05	0.655176

In the next step the SF will be determined via the glassy carbon measurement. This is step (2).

```
In [17]: data GC = np.loadtxt(r'C:\Users\ZechT\FAUbox\Auswertung\Software\ICS
          Py\compute\SAXS\test data/01719 GC.chi')
          sample GCA = np.loadtxt(r'C:\Users\ZechT\FAUbox\Auswertung\Software\
          ICSPy\compute\SAXS\reference/GC APS.txt')
          plt.loglog(data GC[:,0],data GC[:,1],'xg', label='measured GC')
          plt.loglog(sample GCA[:,0], sample GCA[:,1], 'xr', label='reference GC
          1)
          def fit lin(x,a,c):
              return np.exp(a*np.log10(x)+c)
          data BG GC = np.zeros(shape=(len(data GC[:,0])))
          SF, Line GC, Line Ref = calc SF(I GC=data GC[:,1],
                        t GC = 3600,
                        Q GC=data GC[:,0],
                        T GC=table.loc['GC','T'],
                        d GC=0.1)
          d GC=0.1 #cm
          T GC=table.loc['GC','T']
          t GC=3600 #s
          plt.loglog(data_GC[:,0],data_GC[:,1]/(d_GC*t_GC*T_GC),'xb',label='tr
          ansmission corrected GC')
          plt.loglog(data GC[:,0],fit lin(data GC[:,0],Line GC[0],Line GC[1]))
          plt.loglog(sample GCA[:,0],fit lin(sample GCA[:,0],Line Ref[0],Line
          Ref[1]))
          plt.loglog(data_GC[:,0],data_GC[:,1]/(d_GC*t_GC*T_GC)*SF,'xb',label=
          'absolute corrected GC')
          plt.legend(loc='best')
          plt.grid()
          plt.show()
          print('calculated scaling factor SF = ' +str(SF))
          Fit Params. Meas: [0.76870081 0.33778259] Fit Params. Ref:
          74103272 28.43637373]
          84.18543275191664
           10<sup>2</sup>
                          10<sup>1</sup>
           10°
           10^{-1}
                   measured GC
           10^{-2}
                   reference GC
                   transmission corrected GC
           10^{-3}
                   absolute corrected GC
              10-1
                                  10°
                                                      10<sup>1</sup>
          calculated scaling factor SF = 84.18543275191664
```

In step (3) the thickness of the samples will be calculated from H2O and empty cell measurements.

```
In [18]: T_wc = table.loc['H2O','T']
    T_c = table.loc['EC','T']

    d_s = calc_d(T_wc=T_wc,T_c=T_c)
    print('thickness of sample d_s = %f01; (should be about 0.1 cm)' % d_s)

    thickness of sample d_s = 0.09035601; (should be about 0.1 cm)
```

In the end of this procedure (step (4)) the differential scattering cross section of the sample will be calculated, by using the parameters determined in the steps above.

```
In [20]: T_s = table.loc['SGNR','T']
    T_BG = table.loc['H2O','T']

data_s = np.loadtxt(r'C:\Users\ZechT\FAUbox\Auswertung\Software\ICSP
    y\compute\SAXS\test_data/01725_S_GNR.chi')
    data_H2O = np.loadtxt(r'C:\Users\ZechT\FAUbox\Auswertung\Software\IC
    SPy\compute\SAXS\test_data/01745_H2O.chi')

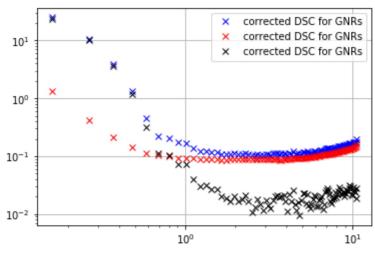
I_s = data_s[:,1]
    BG_s = data_H2O[:,1]

t_s = 3600
    t_BG = 3600

d_BG = float(d_s)

DSC_s = SF*(I_s/(d_s*T_s*t_s)-BG_s/(t_BG*T_BG*d_BG))
```

```
In [23]: plt.loglog(data_s[:,0],data_s[:,1],'xb',label='corrected DSC for GNR
s');
   plt.loglog(data_H2O[:,0],data_H2O[:,1],'xr',label='corrected DSC for
   GNRs');
   plt.loglog(data_s[:,0],DSC_s,'xk',label='corrected DSC for GNRs');
   plt.legend()
   plt.grid()
   plt.show()
```



```
In [ ]:
```