ConventionalPeakFitting

June 18, 2019

```
[1]: import numpy as np
  import matplotlib.pyplot as plt
  from scipy.special import wofz
  from scipy.optimize import curve_fit
  from scipy.optimize import minimize

from tqdm import tqdm_notebook
  from lmfit.models import VoigtModel
  from tomoproc.util.npmath import discrete_cdf
```

The gaussian function is:

$$g(x) = \frac{A}{\sigma\sqrt{2\pi}}exp(\frac{-(x-\mu)^2}{2\sigma^2}),$$

where *A* is the amplitude, μ is the mean (peak center), and σ is the variance.

The half-width at half-maximum (HWHM) for a gaussian distribution, α , can be calculate with

$$\alpha = \sigma \sqrt{2ln(2)}.$$

```
[2]: # functions

def gaussian(x, A, mu, sigma):
    return A/sigma/np.sqrt(2*np.pi)*np.exp(-(x-mu)**2/2/(sigma**2))

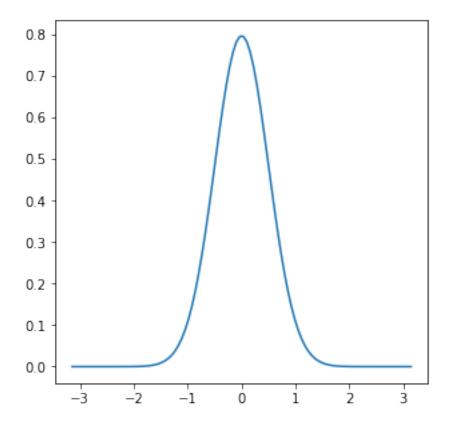
# quick test

x = np.linspace(-np.pi, np.pi, 100)

fig, ax = plt.subplots(1,1,figsize=(5,5))

ax.plot(x, gaussian(x, 1, 0, 0.5))
```

[2]: [<matplotlib.lines.Line2D at 0x1a16e8aba8>]



The Lorentz function (a probability density function, PDF) is:

$$l(x) = \frac{1}{\pi \gamma} \frac{1}{1 + (\frac{x - \mu}{\gamma})^2},$$

where μ is the mean (peak center) and γ is the half-width at half-maximum (HWHM).

Since X-ray diffraction signal from detector is not a proper PDF, the function above need to be adjusted for peak fitting purpose, namely

$$l(x) = I\left[\frac{\gamma^2}{(x-\mu)^2 + \gamma^2}\right],$$

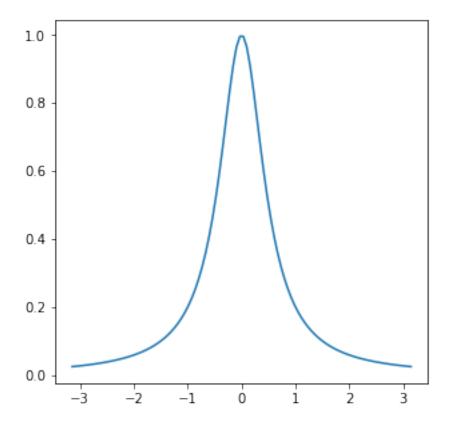
where *I* is the height of the peak.

```
[3]: def lorentz(x, I, mu, gamma):
    return I*(gamma**2)/((x-mu)**2 + gamma**2)

# quick test
x = np.linspace(-np.pi, np.pi, 100)

fig, ax = plt.subplots(1,1,figsize=(5,5))
ax.plot(x, lorentz(x, 1, 0, 0.5))
```

[3]: [<matplotlib.lines.Line2D at 0x1a17227940>]



The Voigt line profile is the convolution of a Gaussian profile, g(x;) and a Lorentzian profile, l(x;),

$$v(x, A, \mu, \sigma, \gamma) = \int_{-\infty}^{+\infty} g(x, 1, \mu, \sigma) \, l(\mu - x, A, \mu, \gamma) dx.$$

```
[4]: def voigt(x, A, mu, sigma, gamma):
    gg = gaussian(x, 1, mu, sigma)
    ll = lorentz(mu-x, A*gamma, mu, gamma)
    return np.convolve(gg, ll, 'same')

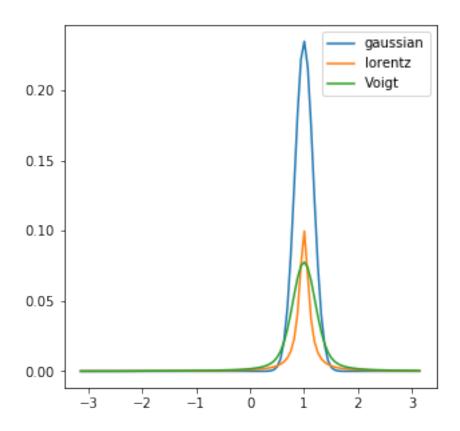
# quick test
x = np.linspace(-np.pi, np.pi, 101)
A = 0.1
mu = 1.0
alpha = 0.2
sigma = alpha/np.sqrt(2*np.log(2))
gamma = 0.1

fig, ax = plt.subplots(1,1,figsize=(5,5))

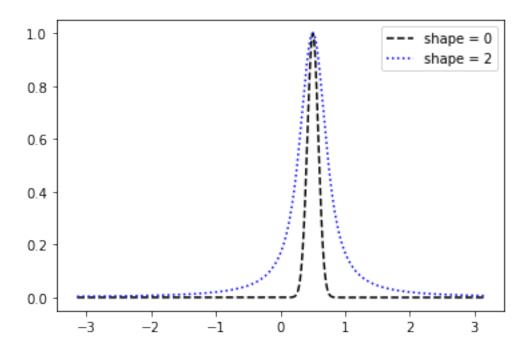
ax.plot(x, gaussian(x, A, mu, sigma), label='gaussian')
ax.plot(x, lorentz(x, A, mu, gamma), label='lorentz')
```

```
ax.plot(x, voigt( x, A, mu, sigma, gamma), label='Voigt')
ax.legend()
```

[4]: <matplotlib.legend.Legend at 0x1a172cf160>



As pointed by this post, convolution is expensive in terms of computation time which can get annoying when used as fit model. The following sample code does not need convolution.

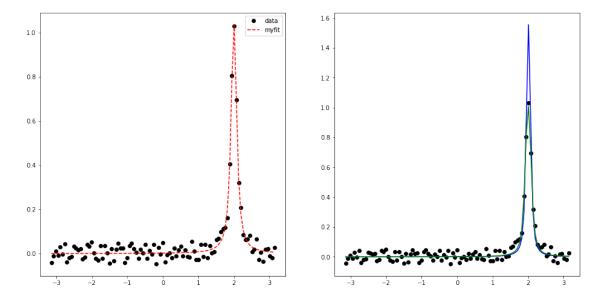


Therefore, we should use the fast version above for peak fitting purpose. Now, let's try to fit some 1D curves.

```
[6]: # generate synthetic signal
    paras = [1, 2, 0.1, 1.5]
    amp, pos, fwhm, shape = paras
    N = 101
    xdata = np.linspace(-np.pi, np.pi, N)
    ydata = voigt(xdata, amp, pos, fwhm, shape) + (np.random.random(N)-0.5)*amp/10
    popt, pcov = curve_fit(voigt, xdata, ydata,
                           maxfev=int(1e6),
                           p0=[ydata.max(), xdata.mean(), 1, 1],
                           bounds=([0,
                                                     xdata.min(), 0,
            0],
                                    [ydata.max()*10, xdata.max(), xdata.max()-xdata.
     →min(), np.inf])
                          )
    fig, ax = plt.subplots(1, 2, figsize=(16, 8))
    ax[0].plot(xdata, ydata, 'ko', label='data')
    ax[0].plot(xdata, voigt(xdata, *popt), 'r--', label='myfit')
    ax[0].legend()
    print(popt)
    from lmfit.models import VoigtModel
```

```
mod = VoigtModel()
pars = mod.guess(ydata, x=xdata)
out = mod.fit(ydata, pars, x=xdata)
ax[1].plot(xdata, ydata, 'ko', label='data')
ax[1].plot(xdata, out.init_fit, 'b-')
ax[1].plot(xdata, out.best_fit, 'g-')
print(out.best_values)
```

```
[1.04187419 2.00290082 0.07106849 2.29185662]
{'sigma': 0.05687740754375503, 'center': 2.0030428821171267, 'amplitude': 0.276136951482719, 'gamma': 0.05687740754375503}
```

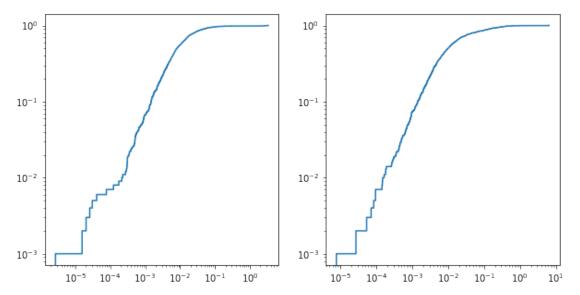


HBox(children=(IntProgress(value=0, max=1000), HTML(value='')))

```
[12]: fig, ax = plt.subplots(1, 2, figsize=(10, 5))

xx, yy = discrete_cdf(np.absolute(dxc_myfit))
ax[0].plot(xx, yy, label='myfit')
ax[0].set_yscale('log')
ax[0].set_xscale('log')

xx, yy = discrete_cdf(np.absolute(dxc_lmfit))
ax[1].plot(xx, yy, label='lmfit')
ax[1].set_yscale('log')
ax[1].set_xscale('log')
```



```
[19]: amp, pos, fwhm, shape = np.random.random(4)*np.pi
    N = 101
    xdata = np.linspace(-np.pi, np.pi, N)
    ydata = voigt(xdata, amp, pos, fwhm, shape) + (np.random.random(N)-0.5)*amp/10

[20]: %timeit x = myfit(xdata, ydata)

14.5 ms ś 609 ţs per loop (mean ś std. dev. of 7 runs, 100 loops each)

[21]: %timeit x = reffit(xdata, ydata)

12.2 ms ś 86.4 ţs per loop (mean ś std. dev. of 7 runs, 100 loops each)

[1]:
```