Cutting pattern

Extracting diffraction orders from Echelle spectrometer spectral image of an incandescent lamp.

imports

```
In [1]:
        %matplotlib inline
In [2]:
        from scipy.signal import savgol_filter
         from matplotlib.pylab import *
        import matplotlib as mpl
        import numpy as np
        import peakutils
```

A typical Echelle spectrometer image with several diffraction orders. Each diffraction order is a band with a parabolic shape. Intensity for

""" Gaussian shape """

return A*np.exp(-((x-c)/s)**2)

A,c,s = arg

Generate test image

each order has a maximum in the center. With increasing diffraction order number, the intensity usually decreases. Spacing between orders, their shape will depend on the set-up. The synthetic image is helpful to demonstrate the extraction procedure of the diffraction orders. In [3]: **def** gf(x, *arg):

```
In [4]:
         # image size in pixels
        nx, ny = [2048, 2048]
        image = np.zeros([nx, ny])
        # generating order pattern
        x = np.arange(nx)
        pt = np.array([(3e-5 * (x - 300 - i ** 2) ** 2 + 120 + i * 193) for i in range(10)]).astype(int)
        # Add Gaussiang profiles along the generated order lines
        width = 200
        profy = np.linspace(0, 2, width)
        # Add intensity attenuation
        intensity = 1 / ((x - nx / 2) ** 2 + 1e6)
        intensity = intensity / intensity.max()
        xx = np.arange(10)
        vv = 1 / ((xx - 0.1) ** 2 + 10)
        vv = vv / vv.max() * 20
        # Add bands
        for j, p in enumerate(pt):
            for i, y in enumerate(p):
                image[y - width // 2 : y + width // 2 + width % 2, i] = gf(profy, vv[j] * intensity[i], 1, 0.3)
         # Add noise
        image += np.random.rand(nx, ny) * 5
        scale = 1
        imshow(image, origin="lower", cmap="binary") # norm=mpl.colors.LogNorm(image.min(), image.max() * scale))
```

```
1000
 750
 500
 250
   0
                  1000
            500
                          1500
                                 2000
Extract peak positions
Check signal, select amplifier
Because intensity for higher orders may drop significantly, for best peak detection it is better to amplify weak intensitys. Here, select this
```

Out[4]:

2000

1750

1500

<matplotlib.image.AxesImage at 0x2568b010700>

amplifier function.

Smooth signal

amp = np.exp(3e-4*x) # amplify weak signal for higher orders

In [5]: x = np.arange(nx)y=image[:,1000] ysm = savgol filter(y, 21, 1)

plot(ysm*amp)

Out[5]:

25

20

plot(y,'k') plot(y*amp)

```
10
         5
                                1000
                                          1500
       Check
In [6]:
         # calculate baseline
         base = peakutils.baseline(ysm,6)
```

Adjust parameters for correct peak detection

1000

[<matplotlib.lines.Line2D at 0x2568fa50ee0>]

[<matplotlib.lines.Line2D at 0x2568e106250>]

ind = peakutils.indexes(ysm-base, thres =.1, min_dist = 50) plot(ysm) plot(base)

10

In [7]:

Out[7]:

In [8]:

```
plot(ind, ysm[ind], 'x')
         [<matplotlib.lines.Line2D at 0x2568f9f9100>]
Out[6]:
         20
         15
```

imshow(image, origin="lower", cmap="binary") # norm=mpl.colors.LogNorm(image.min(), image.max() * scale))

2000 1750

Peak positions over image

x = np.ones(ind.shape[0])*1000

plot(x,ind,'og')

```
1500
 1250
 1000
  750
  500
  250
   0
                          1500
            500
                   1000
                                  2000
Detect all and plot
```

1500

peaks = []pvals = []

peaks.append(ind)

pat_y = np.array(peaks)

Fit quadratics

x = np.arange(nx)

for xind in xselect:

dx = nx//10

n = 4

y=image[:,xind] $ysm = savgol_filter(y, 21, 1)$ base = peakutils.baseline(ysm, 6) ind = peakutils.indexes(ysm-base, thres =.1, min_dist = 180)

print("Shape's good!\nBut check the image, did we get everything?")

fits = [np.poly1d(np.polyfit(xselect,i,2)) for i in pat_y.T]

pattern = np.array([f(x) for f in fits],dtype=int)

Define positions for peak selection. Even distribution from center, which is the brightest.

Adjust peak finding parameters thres and min_dist to get all peaks, and no extra.

xselect = np.arange(nx//2-dx*n,nx//2+dx*n+1, dx,dtype=int)

pvals.append(ysm[ind]) # check if all peaks were detected uniques = np.unique([len(i) for i in peaks]) if uniques.shape[0] > 1: print("Adjust peak finding parameters,", uniques) else:

If not all peaks are detected incorrectly, there will be different number of results for each x pixel in xselec. If so, no fitting is done.

```
# Plot fits
     [plot(p,'C1') for p in pattern]
imshow(image, origin="lower", cmap="binary") # norm=mpl.colors.LogNorm(image.min(), image.max() * scale))
 # Plot peaks
for ind,px in zip(peaks,xselect):
    x = np.ones(ind.shape[0])*px
    plot(x,ind,'oC3')
plot(x,ind,'oC4')
gcf().set_size_inches([8,8])
Shape's good!
But check the image, did we get everything?
2000
1750
1500
1250
1000
 750
 500
```

Save current image and pattern

750

500

250

250

```
In [9]:
         if uniques.shape[0] == 1:
```

1500

1750

2000

np.savetxt('./data/pattern_image_synthetic.txt',image) np.savetxt('./data/pattern.txt',pattern.T,fmt='%.d')

1000

1250