## Creating image masks

When peaks are detected and band shapes fitted properly, each diffraction order now has a vector of coordinates, corresponding to its center. However the band has cirtain width, and it's convinient to define a mask for our image using numpy.

echelle.py uses an array of masks. Each mask shows only one diffraction order.

## ipmorts

```
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
```

## **Load Image and Pattern**

```
In [3]:
         image = np.loadtxt('./data/pattern image synthetic.txt')
         pattern = np.loadtxt('./data/pattern.txt')
In [4]:
         # Show image in log scale
         #imshow(image, origin="lower", cmap="binary",norm=mpl.colors.LogNorm(image.min(), image.max()*1))
         # Show image
         imshow(image, origin="lower", cmap="binary")
        <matplotlib.image.AxesImage at 0x24058a7fdf0>
Out[4]:
        2000
        1750
        1500
        1250
        1000
         750
         500
         250
           0
                         1000
                                       2000
                   500
                                1500
In [5]:
         DIMO = image.shape[0] # DIMension Orders
         DIMW = image.shape[1] # DIMension Wavelength
         dv = 8
         def make mask(ordind, show=False, **kws):
                 converts linear coordinates into 2d mask to mask the image
                 dv = kws.get("dv", 8)
                 l = pattern[:, ordind]
                 cc = np.arange(-dv, dv + 1, 1)
                 ii = ((np.zeros([DIMW, 1]) + cc).T + 1).flatten()
                 jj = np.repeat(np.arange(DIMW)[np.newaxis, ...], dv * 2 + 1, axis=0,).flatten()
                 \verb|mask = (ii.astype(int, copy=False), jj.astype(int, copy=False)||
                 if not show:
                     return mask
                 else:
                     pp = np.zeros((DIMW, DIMO), dtype=bool)
                     pp[mask] = True
                     return pp
         cutting masks = [make mask(i, dv=dv) for i in range(pattern.shape[1])]
```

```
In [6]:
    orderimages = []
    for m in cutting_masks:
        img = image[m]
        order = img.reshape(dv * 2 + 1, DIMW)
        orderimages.append(order)
```

imshow(orderimages[0],aspect=10,cmap='binary')

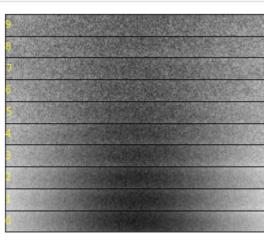
```
0
10 - 250 500 750 1000 1250 1500 1750 2000
```

<matplotlib.image.AxesImage at 0x2405af81c40>

```
In [8]: NORD = pattern.shape[1]
    aspect=10

fig, axs = plt.subplots(NORD, 1)
    plt.subplots_adjust(left=None, bottom=None, right=None, top=None, wspace=0.0, hspace=0.0)
    [ax.set_xticks([]) for ax in axs[:-1]]
    [ax.set_yticks([]) for ax in axs[:-1]]
    [ax.set_xlim([0, DIMW]) for ax in axs]

for o, ax in enumerate(reversed(axs)):
    ax.imshow(orderimages[o], aspect=aspect, cmap='binary')
    ax.get_xaxis().set_visible(False)
    ax.get_yaxis().set_visible(False)
    #ax.axis('off')
    ax.text(
          0, 0.5, o, transform=ax.transAxes, color="#fff83a", va="center", ha="left",
          )
```



## P.S.

Out[7]:

In echelle.py cutting masks are implemented, all we need to supply is the pattern.txt file for the calibration class.

The read\_image method of echelle.py needs to be updated to be independent of image file format. Specify what parameters from the image are used in echelle.py , make a method to read image data and image parameters outside of the class. Probably make a separate python file with read methods.