```
import numpy as np
from astropy.io import fits
import matplotlib.pyplot as plt
from ROHSApy import ROHSA
import scipy.stats as stats
from scipy.signal import find_peaks
from scipy.optimize import curve_fit
import matplotlib.cm as cm
```

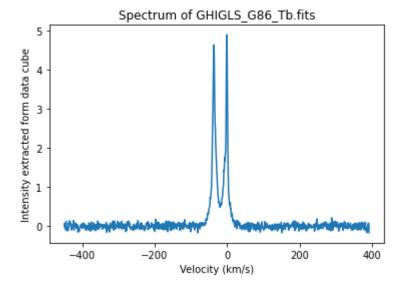
Loading the data cube

```
In [ ]:
    hdul = fits.open("GHIGLS_G86_Tb.fits") # header data unit list
    hdu = hdul[0] #header data unit
    hdr = hdu.header #header
    data_cube = hdu.data[0]
    core = ROHSA(data_cube, hdr, "GHIGLS_G86_Tb.fits")
```

Extracting spectrum for a single pixel

```
pixel_for_plot = (45,45)
spectrum = data_cube[:,pixel_for_plot[0],pixel_for_plot[1]]
velocities = core.v
velocities = np.flip(velocities)
spectrum = np.flip(spectrum)
plt.plot(velocities, spectrum)
plt.xlabel("Velocity (km/s)")
plt.ylabel("Intensity extracted form data cube")
plt.title("Spectrum of GHIGLS_G86_Tb.fits")
```

Out[]: Text(0.5, 1.0, 'Spectrum of GHIGLS_G86_Tb.fits')



~As can be seen above we do have two main peaks!~

Finding Moments of the spectrum

```
def moment(x,p,k,c):
In [ ]:
             Calculates the kth moment of x with distribution p and c as center
             Args:
                 x (np.ndarray): independent variable
                 p (np.ndarray): probability distribution
                 c (float): reference for the moment i.e., k-th moment = sum(p*(x-c)**k)
             return np.sum(((x-c)**k)*p)/np.sum(p)
         # for printing stuff
         def print_moments(velocities, spectrum):
             mean_velocity = moment(velocities, spectrum, 1, 0)
             standard deviation = (moment(velocities, spectrum, 2, mean velocity))**0.5
             skewness = moment(velocities, spectrum, 3, mean_velocity)/standard_deviation**3
             kurtosis = moment(velocities, spectrum, 4, mean_velocity)/standard_deviation**4
             print("Mean velocity:", mean_velocity, "km/s")
             print("Standard deviation:", standard deviation, "km/s")
             print("Skewness:", skewness)
             print("Kurtosis:", kurtosis)
         # Test the moment function
         \# x = np.linspace(0, 10, 100)
         \# p = np.exp(-(x-5)**2/2)
         # plt.plot(x, p)
         # for i in range(1, 3):
               print("Moment", i, ":", moment(x, p, i))
In [ ]:
         # probability distribution from the spectrum
         minimum_possible_intensity = np.min(spectrum)
         prob dist = spectrum - minimum possible intensity
         prob dist = prob dist/np.sum(prob dist)
In [ ]:
         print moments(velocities, spectrum)
         # In units of Column Density
        Mean velocity: -20.432771765296806 km/s
        Standard deviation: 17.102151706247863 km/s
        Skewness: 7.307203281066898
        Kurtosis: -885.6935080484308
```

Dividing the spectrum into two velocity ranges and finding the moments

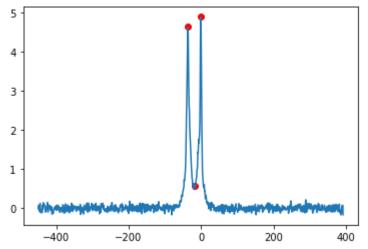
Since there are two peaks we can divide the spectrum into two velocity ranges that contain only a single peak and find the moments. In order to do that we need to find the local minimum of the distribution between the two peaks. And the region that we need to consider is within (-50, 50) km/s since peak are present only in this region.

```
peaks, _ = find_peaks(spectrum, width=1, height=0.5, )
plt.plot(velocities[peaks], spectrum[peaks], "ro")
```

```
plt.plot(velocities, spectrum)
peaks_vel = velocities[peaks]
print("Peaks indexes:", peaks)
print("Peaks are at velocities:", peaks_vel)
print("Initial velocity range:", np.min(velocities), "to", np.max(velocities))
```

Peaks indexes: [514 537 558]

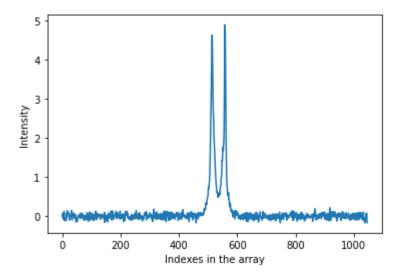
Peaks are at velocities: [-36.19426125 -17.68090832 -0.77541487] Initial velocity range: -449.3317113481132 to 392.6137384951109



The main peaks are at velocities: -36.19426125 km/s and -0.77541487 km/s. These correspond to the indexes 514 and 558 in the spectrum array. And the local minimum is at velocity -17.68090832 km/s. This corresponds to the index 537. So we'll seperate into two velocity ranges: (-100, -17.68090832) km/s and (-17.68090832, 50) km/s. Since the tails are nearly constant apart from the noise so it is not necessary to consider the tails.

```
In [ ]:
    plt.plot(spectrum)
    plt.xlabel("Indexes in the array")
    plt.ylabel("Intensity")
```

Out[]: Text(0, 0.5, 'Intensity')



```
right_part_velocities = velocities[peaks[1]:600]
right_part_spectrum = spectrum[peaks[1]:600]
```

```
left_part_velocities = velocities[400:peaks[1]]
         left part spectrum = spectrum[400:peaks[1]]
In [ ]:
         # Velocitiy Range 1
         plt.title("Left Side Velocity Range: GHIGLS_G86_Tb.fits (-90,-17.68)km/s")
         plt.plot(left_part_velocities, left_part_spectrum)
         print_moments(left_part_velocities, left_part_spectrum)
        Mean velocity: -36.83919964318618 km/s
        Standard deviation: 10.115934829842644 km/s
        Skewness: -2.3645335224101913
        Kurtosis: 14.147285386395446
         Left Side Velocity Range: GHIGLS_G86_Tb.fits (-90,-17.68)km/s
          4
          3
          2
          1
               -120
                       -100
                                -80
                                        -60
                                                -40
                                                        -20
In [ ]:
         # Velocitiy Range 2
         plt.title("Velocity Range 2: GHIGLS_G86_Tb.fits (-17.68, 65)km/s")
         plt.plot(right_part_velocities, right_part_spectrum)
         print_moments(right_part_velocities, right_part_spectrum)
        Mean velocity: -1.5245469218482248 km/s
        Standard deviation: 7.75119627037647 km/s
        Skewness: 0.8754639186230907
        Kurtosis: 5.585697462953123
           Velocity Range 2: GHIGLS_G86_Tb.fits (-17.68, 65)km/s
         5
         4
         3
```

Measuring the noise in the spectrum

10

20

30

2

1

0 -20

-10

Noisy tails are in the range of indexes: (0, 400) and (600, 1047).

```
In []:
    # Left Noisy tail
    left_tail = spectrum[:400]
    avg_noise_left = np.std(left_tail)

# Right Noisy tail
    right_tail = spectrum[600:]
    avg_noise_right = np.std(right_tail)

print("Average noise in the left tail:", avg_noise_left)
    print("Average noise in the right tail:", avg_noise_right)
```

Average noise in the left tail: 0.063227735 Average noise in the right tail: 0.0624478

The noise from the two tails turns out to be nearly the same but with opposite sign. They are very close to 0.

Fitting Gaussians to the spectrum

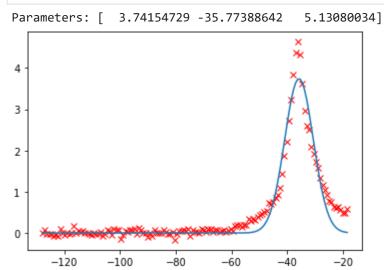
```
def gaussian(x, a, mu, sigma):
    return a*np.exp(-(x-mu)**2/(2*sigma**2))

def gaussian_fit_plot(x, y, initial_guess):
    cf = curve_fit(gaussian, x, y, p0=initial_guess)
    parameters = cf[0]

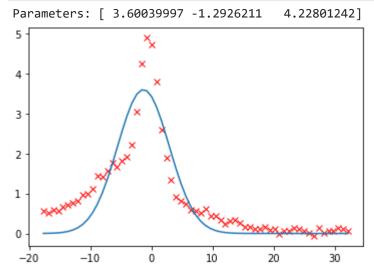
    plt.plot(x, y, "rx")
    plt.plot(x, gaussian(x, *parameters))
    print("Parameters:", parameters)

    return parameters

def gaussian_plot(x, y, params):
    plt.plot(x, y, "rx")
```



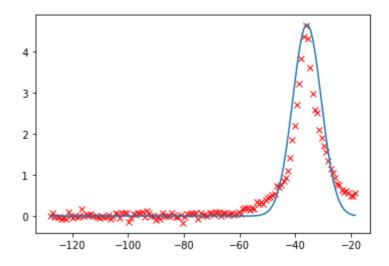




In the above approach we used the initial guess for mean and standard deviation to be the same as the experimentally calculated values. But let's try to find the best fit for the Gaussians by tweaking the parameters a little bit.

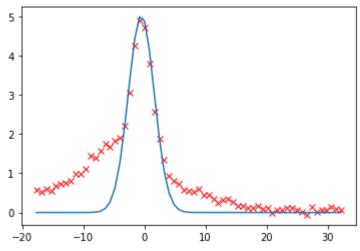
```
# Tweaking Left Part of the spectrum
tweaked_left_params = gaussian_plot(left_part_velocities, left_part_spectrum, [1.25*3.7
```

Parameters: [4.676934112500001, -35.77388642, 5.13080034]

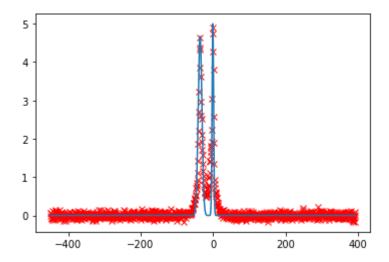


```
# Tweaking Right Part of the spectrum
tweaked_right_params = gaussian_plot(right_part_velocities, right_part_spectrum, [1.4*3]
```

Parameters: [5.040559957999999, -0.49262110000000003, 2.11400621]

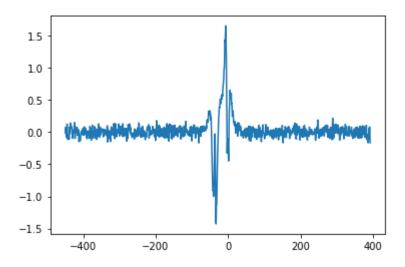


The Combined Gaussian fit



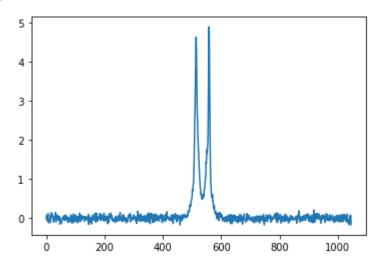
#residual plot
plt.plot(velocities, spectrum - gaussian_combined(velocities, tweaked_left_params, tweaked_left_params)

Out[]: [<matplotlib.lines.Line2D at 0x2065e7df160>]



In []: plt.plot(spectrum)

Out[]: [<matplotlib.lines.Line2D at 0x2065e7e8fd0>]



```
In []: # Cutted Spectrum
print_moments(velocities, spectrum)

Mean velocity: -20.432771765296806 km/s
Standard deviation: 17.102151706247863 km/s
Skewness: 7.307203281066898
Kurtosis: -885.6935080484308

Comparison of the moments
```

1. For the left velocity range

```
In []: # For Left Part of the spectrum
    print("Observed values:")
    print_moments(left_part_velocities, left_part_spectrum)

    print("\nFitted values:")
    print("Mean velocity:", parameters_left[1], "km/s")
    print("Standard deviation:", parameters_left[2], "km/s")

Observed values:
    Mean velocity: -36.83919964318618 km/s
    Standard deviation: 10.115934829842644 km/s
    Skewness: -2.3645335224101913
    Kurtosis: 14.147285386395446

Fitted values:
    Mean velocity: -35.77388641653585 km/s
```

2. For the right velocity range

Standard deviation: 5.1308003351219815 km/s

```
In []: # For Right Part of the spectrum
    print("Observed values:")
    print_moments(right_part_velocities, right_part_spectrum)

    print("\nFitted values:")
    print("Mean velocity:", parameters_right[1], "km/s")

    print("Standard deviation:", parameters_right[2], "km/s")

Observed values:
    Mean velocity: -1.5245469218482248 km/s
    Standard deviation: 7.75119627037647 km/s
    Skewness: 0.8754639186230907
    Kurtosis: 5.585697462953123

Fitted values:
    Mean velocity: -1.2926210950914228 km/s
    Standard deviation: 4.228012420407857 km/s
```

Deriving the Relation between dispersion of the Gaussian and the FWHM of the Gaussian.

For a gaussian of the following form:

$$g(x) = rac{1}{\sqrt{2\pi\sigma^2}} \mathrm{exp}igg(-rac{(x-\mu)^2}{2\sigma^2}igg)$$

The FWHM is the distance between the points along x axis at which the function has a value of half of it's value at the mean. Thus we get

$$rac{1}{2}rac{1}{\sqrt{2\pi\sigma^2}}=rac{1}{\sqrt{2\pi\sigma^2}}\mathrm{exp}igg(-rac{(\mathrm{FWHM}/2)^2}{2\sigma^2}igg)$$
 $rac{1}{2}=\mathrm{exp}igg(-rac{(\mathrm{FWHM}/2)^2}{2\sigma^2}igg)\Rightarrow (2\ln 2)^{1/2}=rac{\mathrm{FWHM}}{2\sigma}$

Thus, we finally get:

$$FWHM = 2 \sigma \sqrt{2 \ln 2} \approx 2.355 \sigma$$

To find FWHM in units of temp. i.e., K,

$$rac{1}{2}mv^2=k_BT$$
 FWHM (in K) $=rac{1}{2k_B}m$ (FWHM (in Km/s)) $^2=rac{4m\ln2}{k_B}\sigma^2$

Here m is the mass of one of the gas particle i.e., aromic hydrogen. Thus $m=1.008*1.67377*10^{-27}kg$. Thus we get:

$$ext{FWHM (in K)} = rac{4m \ln 2}{k_B} \ \sigma^2 pprox 0.00034 \ \sigma^2$$

Here σ^2 is the variance of the Gaussian in unit of (Km/s)^2.

```
In []:
    m = 1.008*1.67377E-27
    k_B = 1.380649E-23

# FWHM for the Left part of the spectrum
    fwhm_left = 2*np.sqrt(2*np.log(2))*parameters_left[2]*1000
    fwhm_left_K = (m/(2*k_B))*((fwhm_left)**2)

# FWHM for the right part of the spectrum
    fwhm_right = 2*np.sqrt(2*np.log(2))*parameters_right[2]*1000
    fwhm_right_K = (m/(2*k_B))*((fwhm_right)**2)

#print("\nFWHM for the Left part of the spectrum:", fwhm_Left, "km/s")
    #print("FWHM for the right part of the spectrum:", fwhm_right, "km/s")

print("\nFWHM (in K) for the left part of the spectrum:", fwhm_left_K, "K")

print("FWHM (in K) for the right part of the spectrum:", fwhm_right_K, "K")
```

FWHM (in K) for the left part of the spectrum: 8919.257662242084 K FWHM (in K) for the right part of the spectrum: 6056.629257580936 K

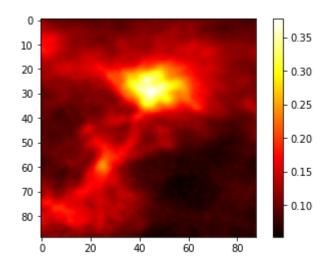
```
In [ ]: ((parameters_left[2]*1E3)**2 )*m/k_B
```

```
Out[]: 3216.9421994318855
```

Avg. 2D map for the whole cube

```
average_spectrum = np.mean(data_cube[:, :,:], axis=0)
    plt.imshow(average_spectrum, cmap=cm.hot)
    plt.colorbar()
```

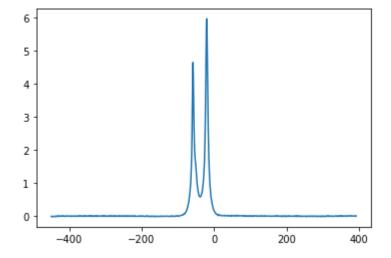
Out[]: <matplotlib.colorbar.Colorbar at 0x2065e6c48e0>



Average spectrum for a small section of the cube

```
In [ ]:
    small_region = data_cube[:, 45-10:45+10, 45-10:45+10]
    avg_spectrum = np.mean(small_region,axis = (1,2))
    plt.plot(velocities, avg_spectrum)
```

Out[]: [<matplotlib.lines.Line2D at 0x2065c5a3c40>]



```
In [ ]:
    # finding the peaks
    peaks_avg, _ = find_peaks(avg_spectrum, width=0.5, height=0.5, )
    plt.plot(velocities[peaks_avg], avg_spectrum[peaks_avg], "ro")
    plt.plot(velocities, avg_spectrum)
```

```
peaks_vel_avg = velocities[peaks_avg]
         print("Peaks indexes:", peaks_avg)
         print("Peaks are at velocities:", peaks_vel_avg)
        Peaks indexes: [487 512 534]
        Peaks are at velocities: [-57.92441016 -37.80400996 -20.09582317]
         5
         3
         2
         1
         0
              -400
                         -200
                                               200
                                                          400
In [ ]:
         plt.plot(avg_spectrum)
         [<matplotlib.lines.Line2D at 0x2065b4222b0>]
Out[]:
         5
         3
         2
         1
         0
                    200
                                      600
                                              800
                             400
                                                      1000
```

Noise in the average spectrum from end-channels

```
In [ ]:
    noise_left_avg = avg_spectrum[:400]
    noise_right_avg = avg_spectrum[600:]

    print("Noise in the left tail:", np.std(noise_left_avg))
    print("Noise in the right tail:", np.std(noise_right_avg))

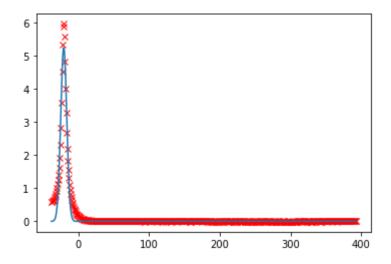
Noise in the left tail: 0.005715796
Noise in the right tail: 0.006848051
```

Gaussian fitting to the average spectrum

```
# Left part of the average spectrum
         left_avg_velocities = velocities[:512]
         left_avg_spectrum = avg_spectrum[:512]
         # Right part of the average spectrum
         right_avg_velocities = velocities[512:]
         right_avg_spectrum = avg_spectrum[512:]
In [ ]:
         print_moments(left_avg_velocities, left_avg_spectrum)
        Mean velocity: -56.475494595667676 km/s
        Standard deviation: 4.931279047848197 km/s
        Skewness: 375.74184419241845
        Kurtosis: -48845.38585375436
In [ ]:
         print_moments(right_avg_velocities, right_avg_spectrum)
        Mean velocity: -19.50592465114145 km/s
        Standard deviation: 10.618807570355564 km/s
        Skewness: 35.97780260306478
        Kurtosis: 2259.8712759742216
In [ ]:
         left_avg_guesses = [np.max(left_avg_spectrum), -56.475, 4.931]
         right_avg_guesses = [np.max(right_avg_spectrum), -19.506, 10.619]
         left avg params = gaussian fit plot(left avg velocities, left avg spectrum, left avg gu
        Parameters: [ 3.45756812 -57.21576853
                                                  5.29100578]
         4
        3
        2
        1
        0
                           -300
                                     -200
                -400
                                                -100
```

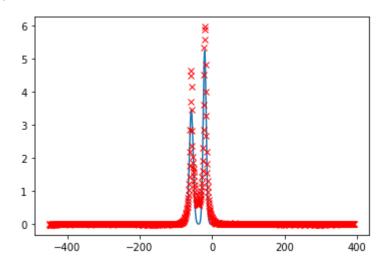
```
In [ ]: right_avg_params = gaussian_fit_plot(right_avg_velocities, right_avg_spectrum, right_avg_spectrum)
```

Parameters: [5.26204505 -20.29384833 4.24610697]



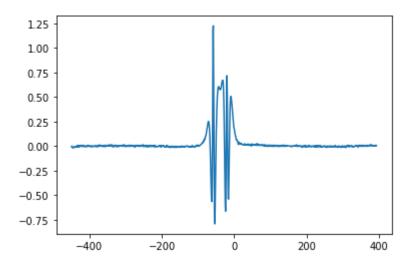
gaussian_combined_avg = gaussian_combined(velocities, left_avg_params, right_avg_params
plt.plot(velocities, gaussian_combined_avg)
plt.plot(velocities, avg_spectrum, "rx")

Out[]: [<matplotlib.lines.Line2D at 0x2065e8ad910>]



Residual Plot
plt.plot(velocities, avg_spectrum - gaussian_combined_avg)

Out[]: [<matplotlib.lines.Line2D at 0x2065e912d00>]



Using ROHSA to decompose the average spectrum

```
In [ ]:
         filename = "GHIGLS G86 Tb.dat" # @param
         fileout = "GHIGLS_G86_Tb_gauss_run_0.dat" #@param
         filename noise = ''
         n_{gauss} = 3
                               #@param {type:"slider", min:1, max:12, step:1}
                               #@param {type:"slider", min:0, max:1000, step:1}
         lambda amp = 100
                               #@param {type:"slider", min:0, max:1000, step:1}
         lambda mu = 100
                               #@param {type:"slider", min:0, max:1000, step:1}
         lambda_sig = 100
                               #@param {type:"slider", min:0, max:10000, step:1}
         lambda_var_sig = 0.
         amp_fact_init = 0.66 # times max amplitude of Gaussians for the fit of the mean spectr
         sig init = 4.
                               # dispersion of Gaussians for the fit of the mean spectrum
                               # lower limit on sigma for the fit of the mean spectrum
         lb_sig_init = 1.
         ub_sig_init = 12.
                               # upper limit on sigma for the fit of the mean spectrum
         lb\_sig = 1.
         ub sig = 100.
         maxiter_init = 15000 # max iteration for L-BFGS-B alogorithm init mean
         maxiter = 800
                               #@param {type:"slider", min:1, max:800, step:1}
         noise = ".false."
                               # if false - STD map computed by ROHSA between lstd and ustd
                               #@param {type:"slider", min:1, max:400, step:1}
         lstd = 1
                               #@param {type:"slider", min:1, max:400, step:1}
         ustd = 20
         iprint = -1
                               #@param ["-1", "0", "1"]
                               # print option init
         iprint init = -1
         save grid = ".false." #@param [".true.", ".false."]
         core = ROHSA(small_region[400:600, :, :], hdr, filename="GHIGLS_G86_Tb.fits")
         core.cube2dat(filename=filename)
         core.gen_parameters(filename=filename,
                             fileout=fileout,
                             n gauss=n gauss,
                             lambda_amp=lambda_amp,
                             lambda_mu=lambda_mu,
                             lambda_sig=lambda_sig,
                             lambda var sig=lambda var sig,
                             amp fact init=amp fact init,
                             sig init=sig init,
                             lb_sig_init=lb_sig_init,
                             ub_sig_init=ub_sig_init,
                             lb_sig=lb_sig,
                             ub_sig=ub_sig,
                             noise=noise,
                             maxiter=maxiter,
```

```
lstd=lstd,
ustd=ustd,
iprint_init=iprint_init,
iprint=iprint,
save_grid=save_grid)
```

Generate GHIGLS_G86_Tb.dat file readable by fortran Generate parameters.txt file

```
def multiple_gaussian(x, parameters):
    # parameters = [amp, mu, sigma, amp, mu, sigma, ...]
    y = 0
    for a,mu,sig in zip(parameters[::3], parameters[1::3], parameters[2::3]):
        y += a*np.exp(-(1/2)*((x-mu)/sig)**2)
    return y
```

Using 3 gaussians we get the following parameters:

```
In []:
    gaussian_from_ROHSA = core.read_gaussian("GHIGLS_G86_Tb_gauss_run_0.dat")
    model_from_ROHSA = core.return_result_cube(gaussian_from_ROHSA)

amplitudes = gaussian_from_ROHSA[0::3]
    positions = gaussian_from_ROHSA[1::3]
    dispersions = gaussian_from_ROHSA[2::3]

actual = small_region[400:600, :, :]

actual_data = plt.plot(np.mean(actual, axis=(1,2)), label="Actual Data")
    model = plt.plot(np.mean(model_from_ROHSA, axis=(1,2)), label="ROHSA Model")

plt.title("GHIGLS G86 Tb: ROHSA Model vs Average spectrum")
    plt.legend()
```

Opening data file <matplotlib.legend.Legend at 0x2065e9366d0>

Out[]:

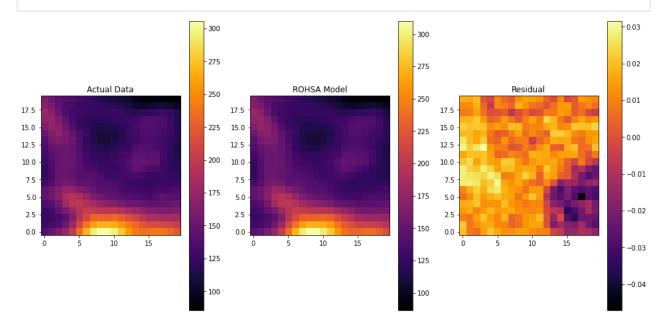
GHIGLS G86 Tb: ROHSA Model vs Average spectrum 6 Actual Data ROHSA Model 5 4 3 2 1 0 125 175 25 50 75 100 150 200

Integrated Column Density Maps

```
def plot_integrated_column_density(cube, rohsa_model):
    plt.figure(figsize=(16, 8))
```

```
plt.subplot(1, 3, 1)
    plt.imshow(np.sum(cube, 0), origin="lower", cmap="inferno")
    plt.title("Actual Data")
    plt.colorbar()
    plt.subplot(1, 3, 2)
    plt.imshow(np.sum(rohsa_model, 0), origin="lower", cmap="inferno")
    plt.title("ROHSA Model")
    plt.colorbar()
    plt.subplot(1, 3, 3)
    plt.imshow((np.sum(rohsa_model, 0)-np.sum(cube, 0)) /
            np.sum(rohsa_model, 0), origin="lower", cmap="inferno")
    plt.title("Residual")
    plt.colorbar()
def plot gaussian parameters(amplitude, position, dispersion):
    plt.figure(figsize=(16, 8))
    plt.subplot(1, 3, 1)
    plt.imshow(amplitude, origin="lower", cmap="inferno")
    plt.title("Amplitudes")
    plt.colorbar()
    plt.subplot(1, 3, 2)
    plt.imshow(position, origin="lower", cmap="coolwarm")
    plt.title("Mean positions")
    plt.colorbar()
    plt.subplot(1, 3, 3)
    plt.imshow(dispersion, origin="lower", cmap="cubehelix")
    plt.title("Dispersions")
    plt.colorbar()
def plot_mosaic_spectra(cube_,integral,rohsa_model,gaussian_from_rohsa):
    #Plot mosaic spectra
    pvalues = np.logspace(-1, 0, len(integral))
    pmin = pvalues[0]
    pmax = pvalues[-1]
    def norm(pval):
        return (pval - pmin) / float(pmax - pmin)
    ny = 4
    nx = 4
    center_y = int(cube_.shape[2]/2)
    center x = int(cube .shape[1]/2)
    x = np.arange(cube .shape[0])
    cb = "magenta"
    cw = "crimson"
    fig, axs = plt.subplots(4, 4, sharex=True, sharey=True, figsize=(10., 6.))
    fig.subplots adjust(hspace=0, wspace=0, left=0, right=1, top=1, bottom=0)
    for i in np.arange(ny):
        for j in np.arange(nx):
            axs[i][j].step(x, cube_[:, center_y+i, center_x+j],
                        color='cornflowerblue', linewidth=2.)
```

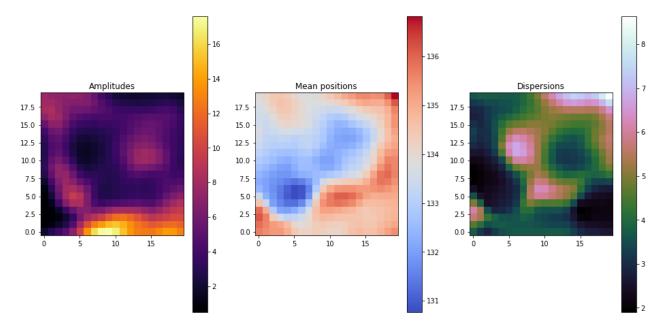
In []: plot_integrated_column_density(actual, model_from_ROHSA)



Map of amplitudes, centers and widths of the Gaussian fits to the average spectrum.

For the first Gaussian:

```
In [ ]: plot_gaussian_parameters(amplitudes[0], positions[0], dispersions[0])
```



Plotting mosaic spectra

In []: plot_mosaic_spectra(actual, amplitudes*dispersions, model_from_ROHSA, gaussian_from_ROH

