

```
In [ ]: 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
```

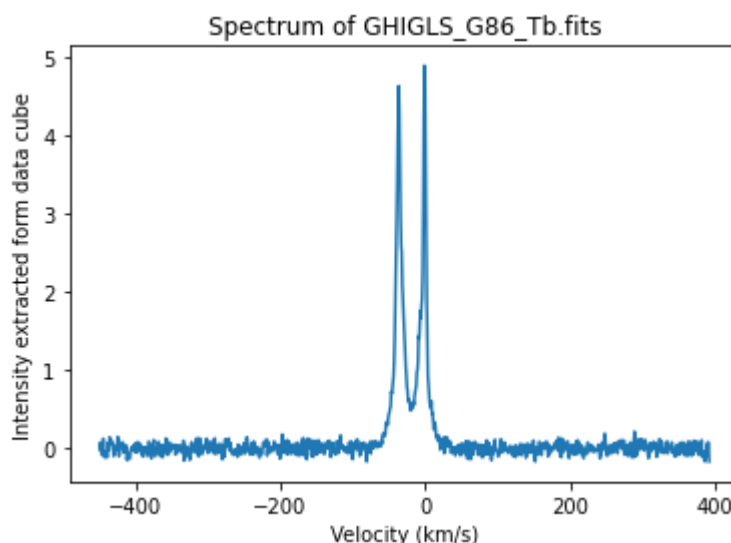
Loading the data cube

```
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

```
In [ ]: 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

```
In [ ]: def moment(x,p,k,c):
        """
        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.

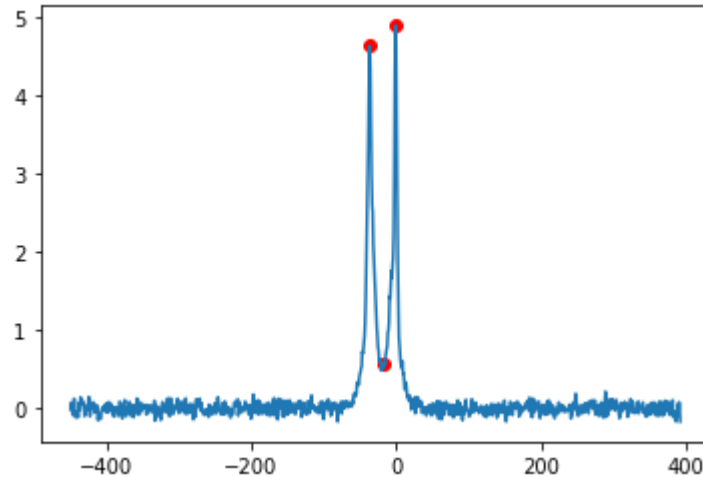
```
In [ ]: peaks, _ = find_peaks(spectrum, width=1, height=0.5, )
        plt.plot(velocities[peaks], spectrum[peaks], "ro")
```

```
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]

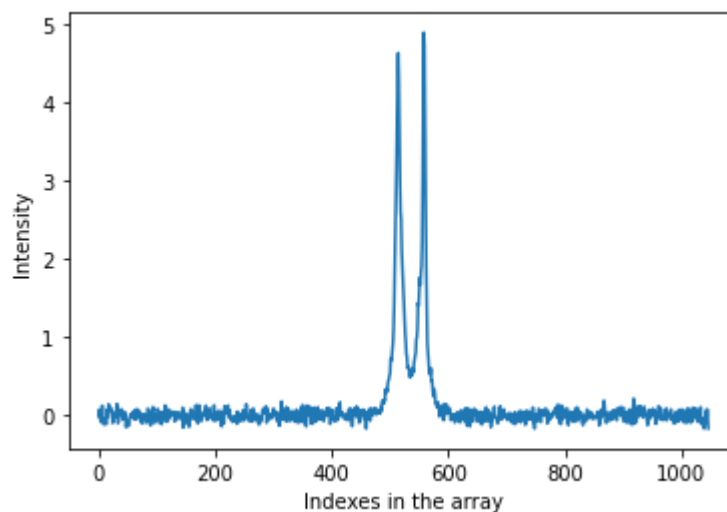
Initial velocity range: -449.3317113481132 to 392.6137384951109



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 separate 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')
```



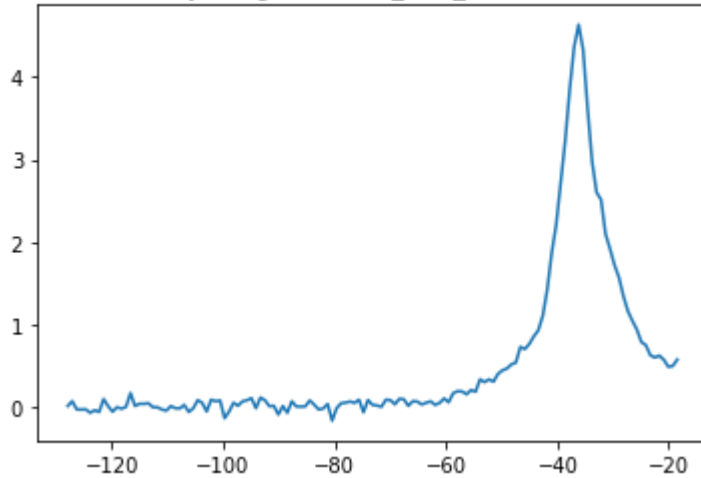
```
In [ ]: right_part_velocities = velocities[peaks[1]:600]
right_part_spectrum = spectrum[peaks[1]:600]
```

```
left_part_velocities = velocities[400:peaks[1]]
left_part_spectrum = spectrum[400:peaks[1]]
```

```
In [ ]: # Velocity 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

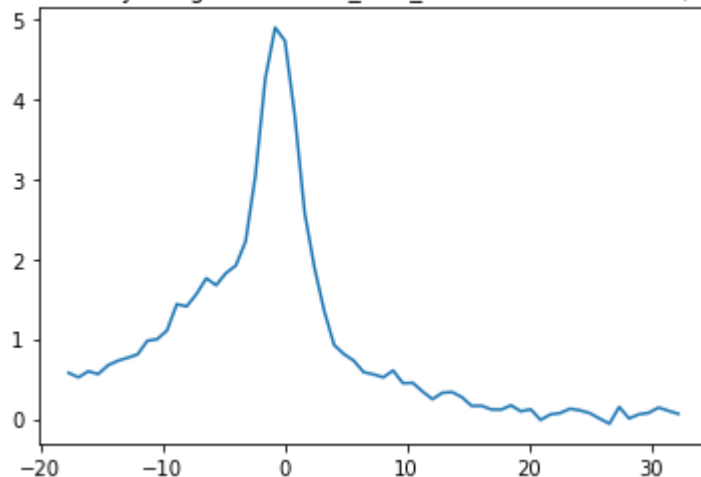
Left Side Velocity Range: GHIGLS_G86_Tb.fits (-90,-17.68)km/s



```
In [ ]: # Velocity 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

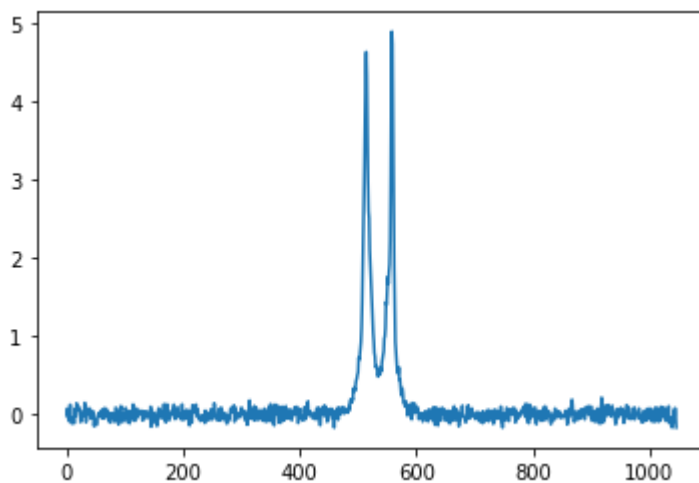
Velocity Range 2: GHIGLS_G86_Tb.fits (-17.68, 65)km/s



Measuring the noise in the spectrum

```
In [ ]: plt.plot(spectrum)
```

```
Out[ ]: [matplotlib.lines.Line2D at 0x2065ea31940]
```



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

```
In [ ]: 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)

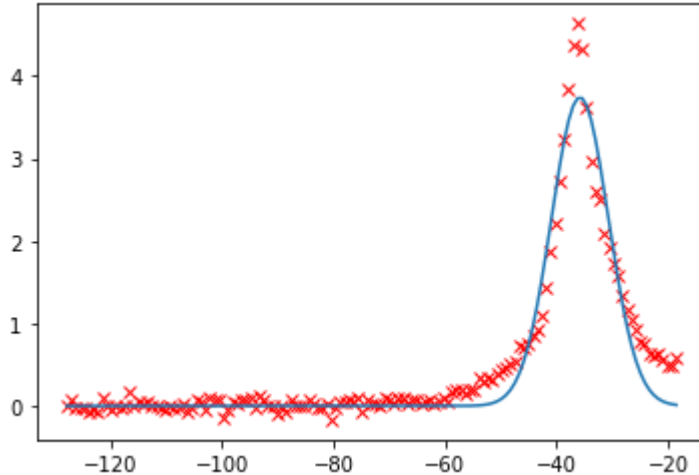
    return parameters

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

```
plt.plot(x, gaussian(x, *params))
print("Parameters:", params)
return params

# Left Part of the spectrum
parameters_left = gaussian_fit_plot(left_part_velocities, left_part_spectrum,
                                    [np.max(left_part_spectrum), -36.8, 102])
```

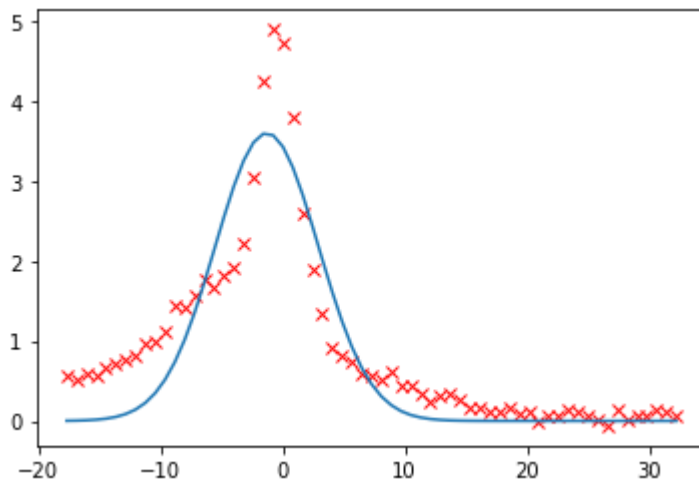
Parameters: [3.74154729 -35.77388642 5.13080034]



In []:

```
# Right Part of the spectrum
parameters_right = gaussian_fit_plot(right_part_velocities, right_part_spectrum, [np.ma
```

Parameters: [3.60039997 -1.2926211 4.22801242]

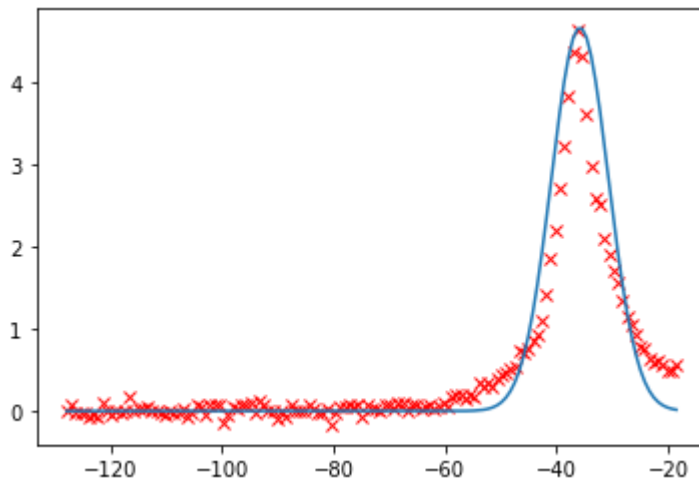


In the above approach we used the initial guess for mean and standard deviation to be the same as the experimentally calculated values. But let's try to find the best fit for the Gaussians by tweaking the parameters a little bit.

In []:

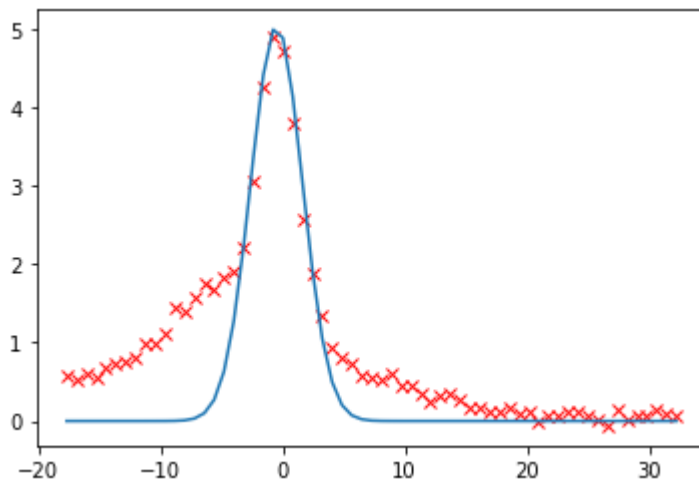
```
# Tweaking Left Part of the spectrum
tweaked_left_params = gaussian_plot(left_part_velocities, left_part_spectrum, [1.25*3.7
```

Parameters: [4.676934112500001, -35.77388642, 5.13080034]



```
In [ ]: # 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]

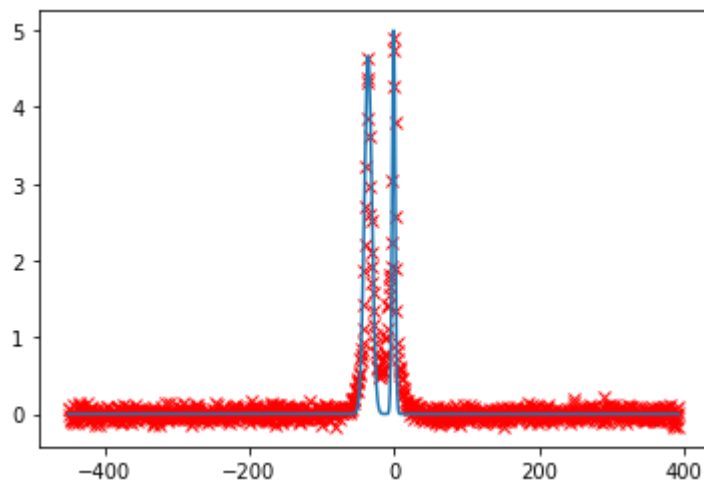


The Combined Gaussian fit

```
In [ ]: def gaussian_combined(x, param_left, param_right):
        return gaussian(x, *param_left) + gaussian(x, *param_right)

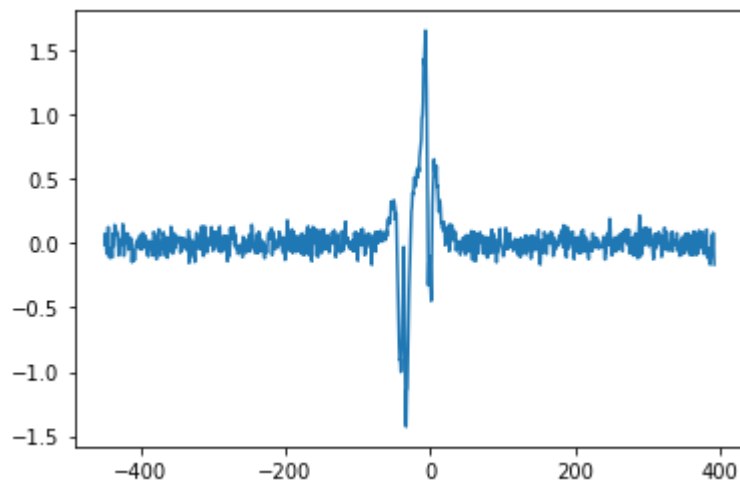
plt.plot(velocities, spectrum, "rx")
plt.plot(velocities, gaussian_combined(velocities, tweaked_left_params, tweaked_right_p
```

Out[]: [matplotlib.lines.Line2D at 0x2065eb5bcd0]



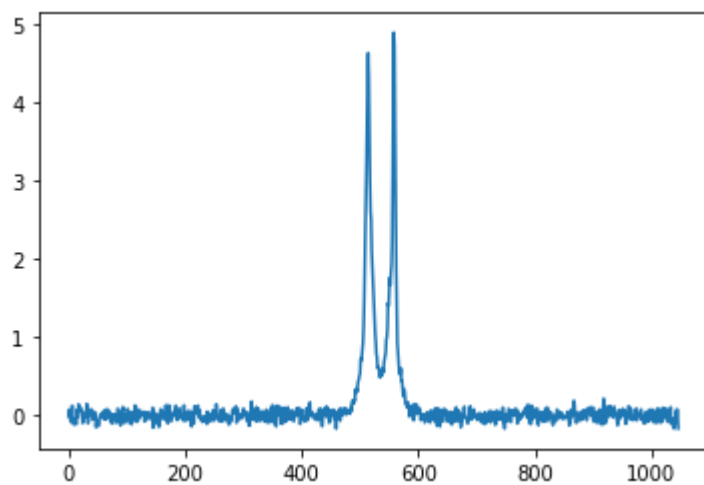
```
In [ ]: #residual plot
plt.plot(velocities, spectrum - gaussian_combined(velocities, tweaked_left_params, twea
```

```
Out[ ]: [matplotlib.lines.Line2D at 0x2065edddf70]
```



```
In [ ]: plt.plot(spectrum)
```

```
Out[ ]: [matplotlib.lines.Line2D at 0x2065eddc1f0]
```




```
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
Standard deviation: 5.1308003351219815 km/s

2. For the right velocity range

```
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:

$$g(x) = \frac{1}{\sqrt{2\pi\sigma^2}} \exp\left(-\frac{(x-\mu)^2}{2\sigma^2}\right)$$

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

$$\begin{aligned} \frac{1}{2} \frac{1}{\sqrt{2\pi\sigma^2}} &= \frac{1}{\sqrt{2\pi\sigma^2}} \exp\left(-\frac{(\text{FWHM}/2)^2}{2\sigma^2}\right) \\ \frac{1}{2} &= \exp\left(-\frac{(\text{FWHM}/2)^2}{2\sigma^2}\right) \Rightarrow (2\ln 2)^{1/2} = \frac{\text{FWHM}}{2\sigma} \end{aligned}$$

Thus, we finally get:

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

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

$$\frac{1}{2}mv^2 = k_B T$$

$$\text{FWHM (in K)} = \frac{1}{2k_B} m (\text{FWHM (in Km/s)})^2 = \frac{4m \ln 2}{k_B} \sigma^2$$

Here m is the mass of one of the gas particle i.e., atomic hydrogen. Thus

$m = 1.008 * 1.67377 * 10^{-27} \text{ kg}$. Thus we get:

$$\text{FWHM (in K)} = \frac{4m \ln 2}{k_B} \sigma^2 \approx 0.00034 \sigma^2$$

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
```

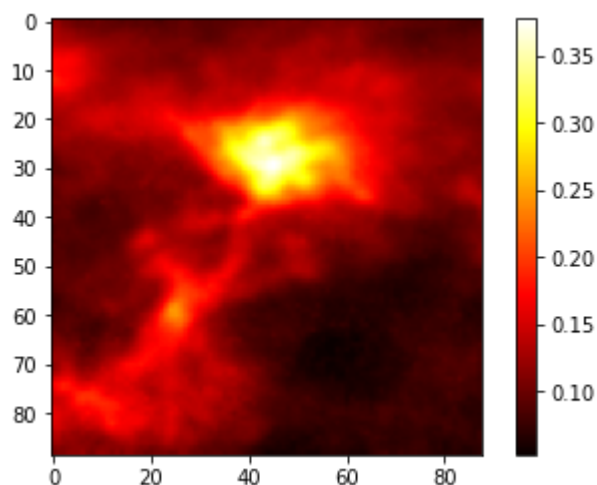
```
In [ ]: ((parameters_left[2]*1E3)**2)*m/k_B
```

Out[]: 3216.9421994318855

Avg. Column density for the whole cube

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

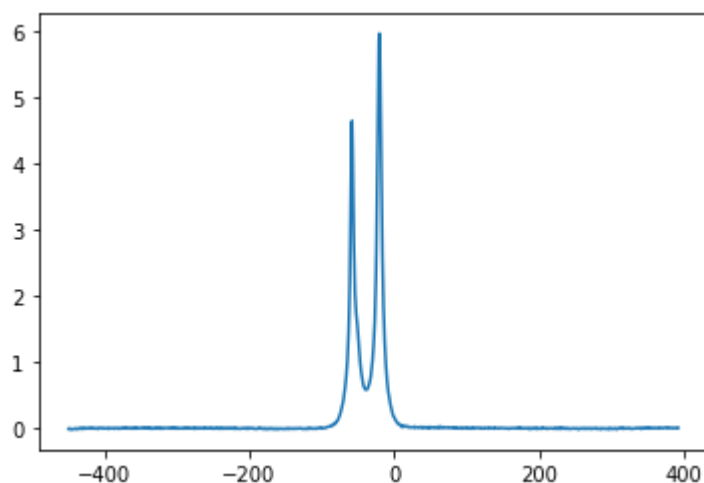
Out[]: <matplotlib.colorbar.Colorbar at 0x2065ec79e80>



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 0x2065eaf97f0>]

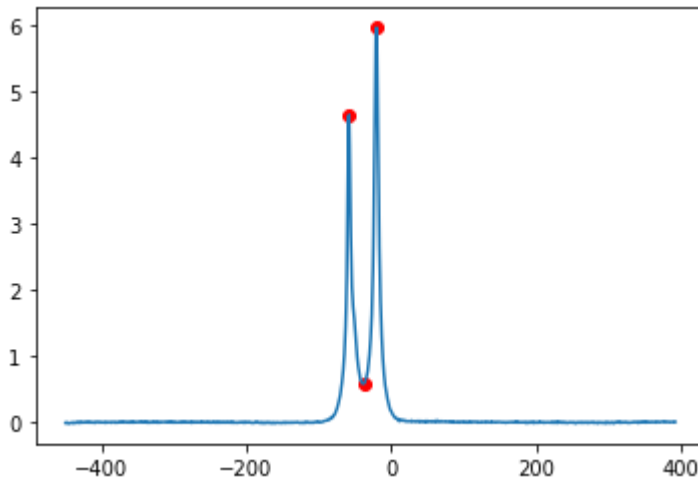


```
In [ ]: # finding the peaks
peaks_avg, _ = find_peaks(avg_spectrum, width=0.5, height=0.5, )
plt.plot(velocities[peaks_avg], avg_spectrum[peaks_avg], "ro")
plt.plot(velocities, avg_spectrum)
```

```
peaks_vel_avg = velocities[peaks_avg]
print("Peaks indexes:", peaks_avg)
print("Peaks are at velocities:", peaks_vel_avg)
```

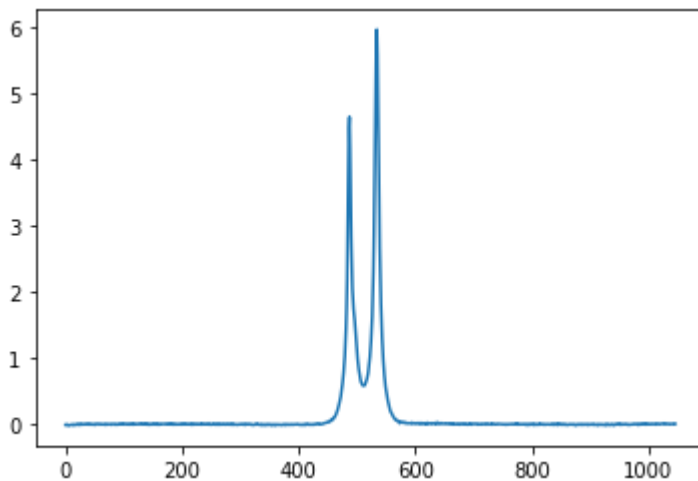
Peaks indexes: [487 512 534]

Peaks are at velocities: [-57.92441016 -37.80400996 -20.09582317]



```
In [ ]: plt.plot(avg_spectrum)
```

```
Out[ ]: [<matplotlib.lines.Line2D at 0x2065eedf850>]
```



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

```
In [ ]:
```

```
# 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
```

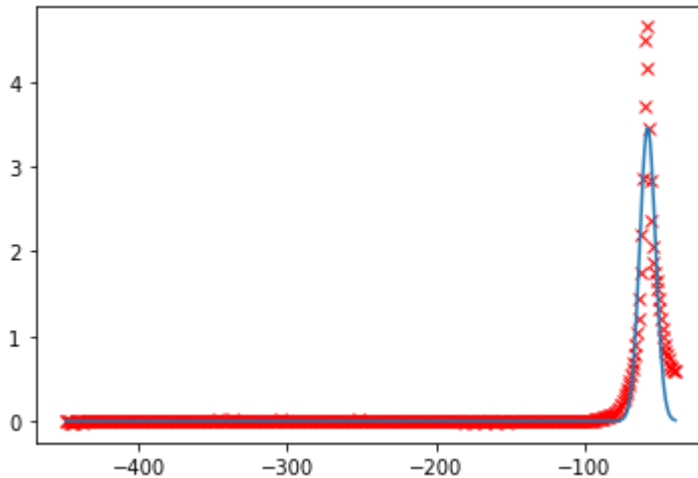
```
In [ ]: print_moments(right_avg_velocities, right_avg_spectrum)
```

```
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]

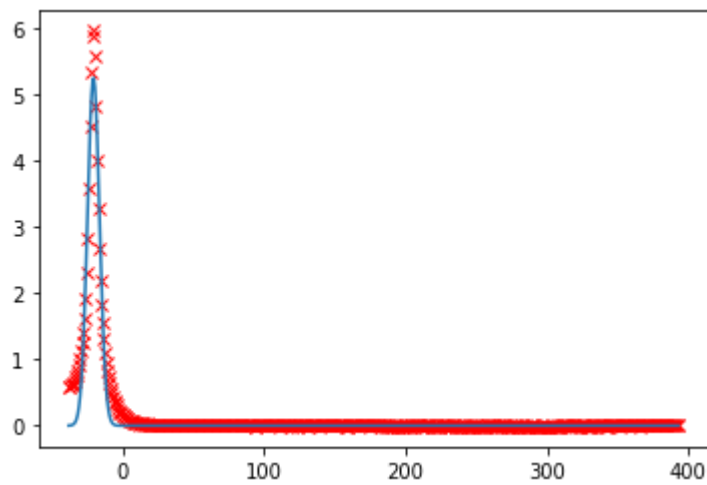
left_avg_params = gaussian_fit_plot(left_avg_velocities, left_avg_spectrum, left_avg_gu
```

```
Parameters: [ 3.45756812 -57.21576853  5.29100578]
```



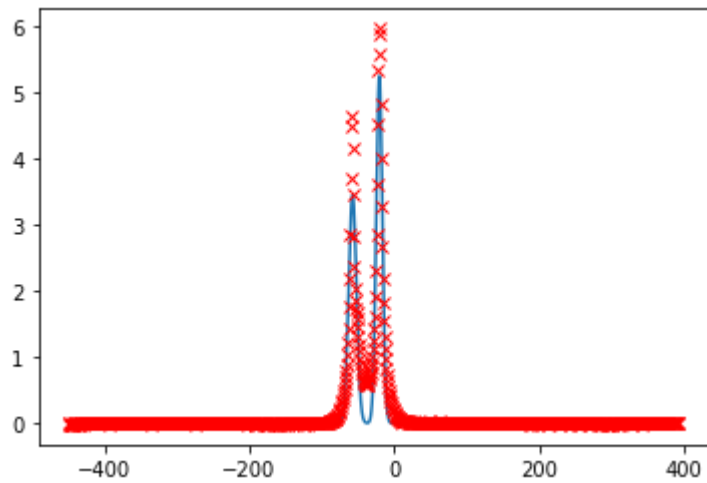
```
In [ ]: right_avg_params = gaussian_fit_plot(right_avg_velocities, right_avg_spectrum, right_av
```

```
Parameters: [ 5.26204505 -20.29384833  4.24610697]
```



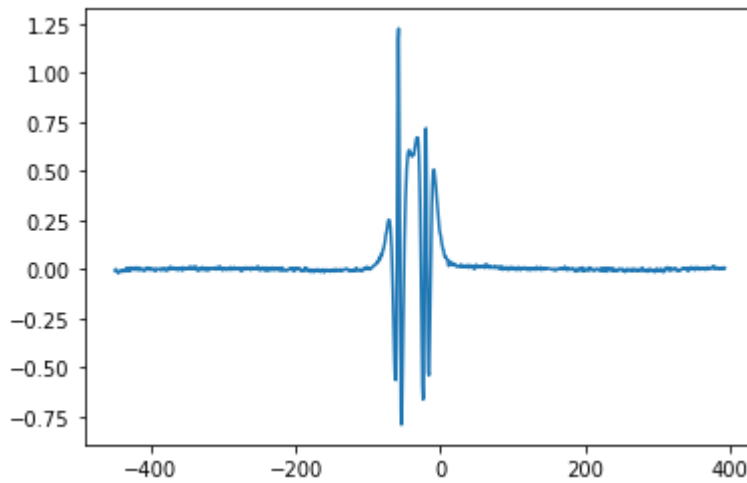
```
In [ ]: gaussian_combined_avg = gaussian_combined(velocities, left_avg_params, right_avg_params)
plt.plot(velocities, gaussian_combined_avg)
plt.plot(velocities, avg_spectrum, "rx")
```

```
Out[ ]: [<matplotlib.lines.Line2D at 0x2065c4bb760>]
```



```
In [ ]: # Residual Plot
plt.plot(velocities, avg_spectrum - gaussian_combined_avg)
```

```
Out[ ]: [<matplotlib.lines.Line2D at 0x2065ee21bb0>]
```



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}
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:1000, step:1}
lambda_var_sig = 0. # @param {type:"slider", min:0, max:10000, step:1}
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
lb_sig_init = 1. # lower limit on sigma for the fit of the mean spectrum
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
lstd = 1 # @param {type:"slider", min:1, max:400, step:1}
ustd = 20 # @param {type:"slider", min:1, max:400, step:1}
iprint = -1 # @param ["-1", "0", "1"]
iprint_init = -1 # print option init
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

```

In [ ]: 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()

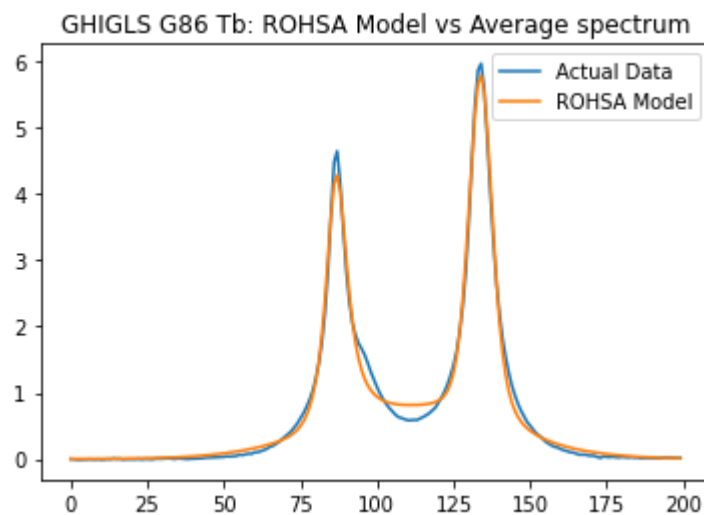
```

Opening data file

```

Out[ ]: <matplotlib.legend.Legend at 0x2065f033070>

```



Integrated Column Density Maps

```

In [ ]: 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.)

```

```

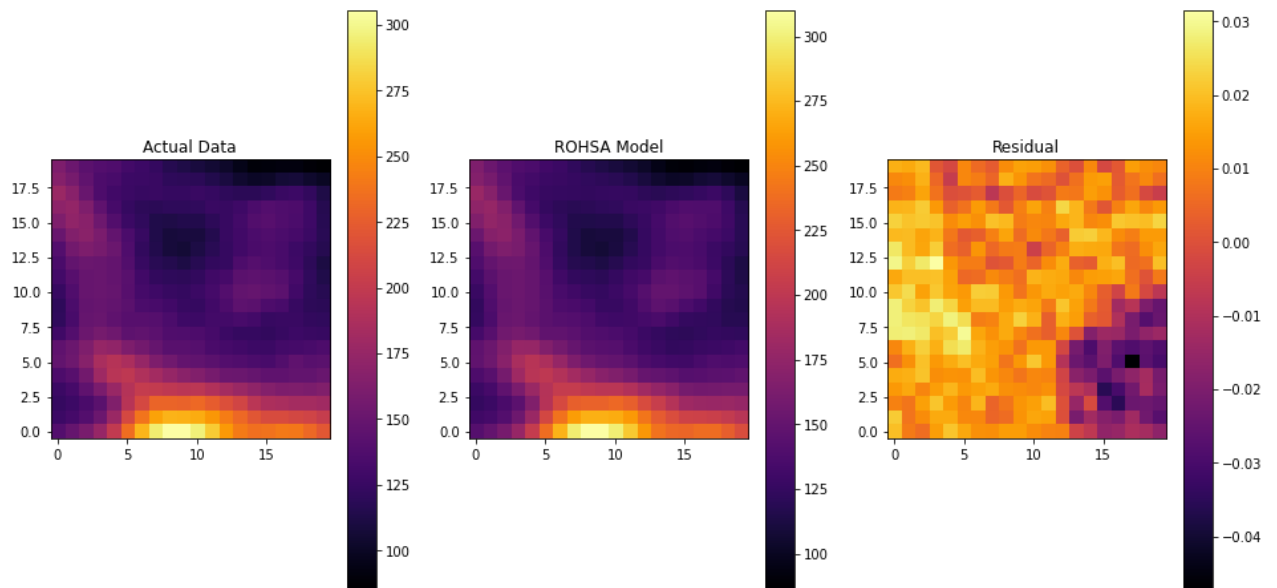
    axs[i][j].plot(x, rohsa_model[:, center_y+i, center_x+j],
                   linestyle="-", linewidth=2., color="k")
    for k in range(len(integral)):
        axs[i][j].plot(x, core.gauss(x, gaussian_from_rohsa[0:3][k][center_y+i,
        gaussian_from_rohsa[1:3][k][center_y +
        i, center_x+j],
        gaussian_from_rohsa[2:3][k][center_y+i, ce
        linewidth=2., color=plt.cm.inferno(pvalues[k]))
    if j == 0:
        axs[i][j].set_ylabel(r'T [k]')
    axs[i][j].set_xlabel(r'v [km s-1 $]')

```

```

In [ ]: plot_integrated_column_density(actual, model_from_ROHSA)

```



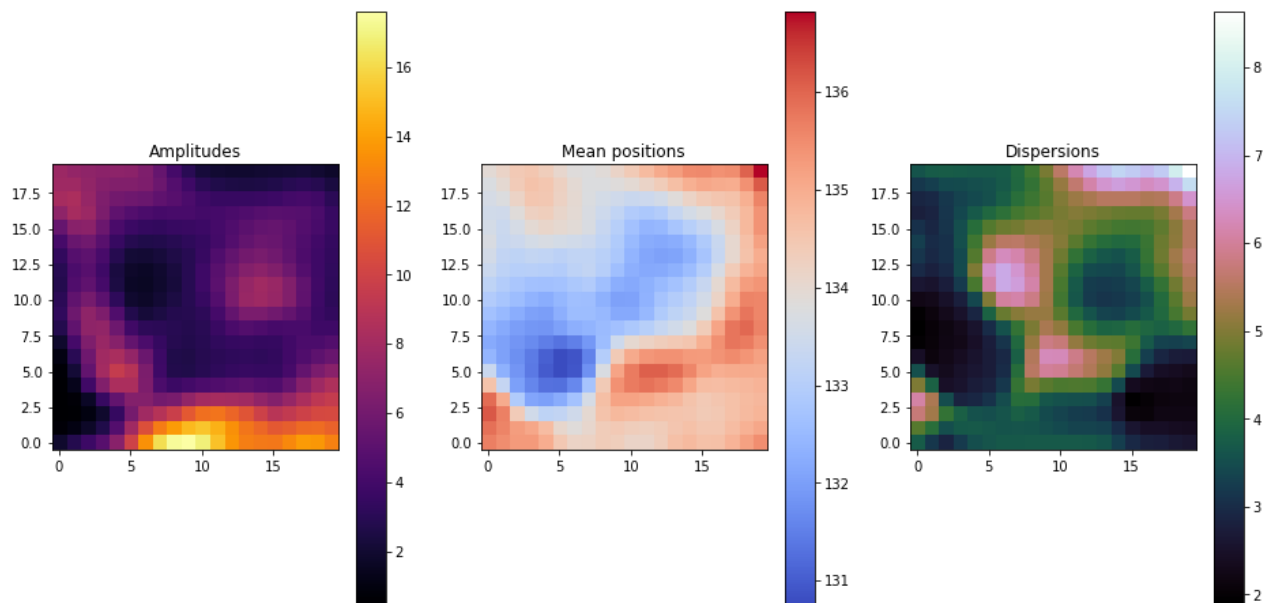
Map of amplitudes, centers and widths of the Gaussian fits to the average spectrum.

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
```

