

# H2D2 Spectroscopy

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```

In [23]: import numpy as np
import pylab as py
import matplotlib.pyplot as plt
from scipy.optimize import curve_fit
import scipy.stats as stats

def fgaussian(x, A, B, C, D):
    return A * np.exp(-((x - B) ** 2) / (2 * C ** 2)) + D

def ftwogaussian(x, A1, A2, B1, B2, C1, C2, D):
    return (A1 * np.exp(-((x - B1) ** 2) / (2 * C1 ** 2))
            + A2 * np.exp(-((x - B2) ** 2) / (2 * C2 ** 2))) + D

def is_float(string):
    try:
        float(string)
        return True
    except ValueError:
        return False

# pull data
#data1 = np.genfromtxt('MercuryData.csv', delimiter=',', skip_header=22, dtype=st
datam = np.genfromtxt('DataFiles/MercuryData.csv', delimiter=',', dtype=st
datam_copy = np.genfromtxt('DataFiles/MercuryDataCopy.csv', delimiter=',',

data1 = np.genfromtxt('DataFiles/H2D2Data1.csv', delimiter=',', dtype=str
data1_peak1 = np.genfromtxt('DataFiles/H2D2Data1Peak1.csv', delimiter=',',
data1_peak2 = np.genfromtxt('DataFiles/H2D2Data1Peak2.csv', delimiter=',',

data2 = np.genfromtxt('DataFiles/H2D2Data2.csv', delimiter=',', dtype=str
data2_peak1 = np.genfromtxt('DataFiles/H2D2Data2Peak1.csv', delimiter=',',
data2_peak2 = np.genfromtxt('DataFiles/H2D2Data2Peak2.csv', delimiter=',',

data3 = np.genfromtxt('DataFiles/H2D2Data3.csv', delimiter=',', dtype=str
data3_peak1 = np.genfromtxt('DataFiles/H2D2Data3Peak1.csv', delimiter=',',
data3_peak2 = np.genfromtxt('DataFiles/H2D2Data3Peak2.csv', delimiter=',',

data4 = np.genfromtxt('DataFiles/H2D2Data4.csv', delimiter=',', dtype=str
data4_peak1 = np.genfromtxt('DataFiles/H2D2Data4Peak1.csv', delimiter=',',
data4_peak2 = np.genfromtxt('DataFiles/H2D2Data4Peak2.csv', delimiter=',',

data5 = np.genfromtxt('DataFiles/Full_Data_copy.csv', delimiter=',', dtype=

# split columns from data into x and y values
x_data_m = [float(row[0]) if is_float(row[0]) else np.nan for row in datam]
y_data_m = [float(row[1]) if is_float(row[1]) else np.nan for row in datam]
x_data_m_Peak1 = [float(row[0]) if is_float(row[0]) else np.nan for row in
y_data_m_Peak1 = [float(row[1]) if is_float(row[1]) else np.nan for row in

x_data_1 = [float(row[0]) if is_float(row[0]) else np.nan for row in data1]
y_data_1 = [float(row[1]) if is_float(row[1]) else np.nan for row in data1]
x_data_1_peak1 = [float(row[0]) if is_float(row[0]) else np.nan for row in
y_data_1_peak1 = [float(row[1]) if is_float(row[1]) else np.nan for row in
x_data_1_peak2 = [float(row[0]) if is_float(row[0]) else np.nan for row in
y_data_1_peak2 = [float(row[1]) if is_float(row[1]) else np.nan for row in

x_data_2 = [float(row[0]) if is_float(row[0]) else np.nan for row in data2]

```

```
y_data_2 = [float(row[1]) if is_float(row[1]) else np.nan for row in data2]
x_data_2_peak1 = [float(row[0]) if is_float(row[0]) else np.nan for row in data2]
y_data_2_peak1 = [float(row[1]) if is_float(row[1]) else np.nan for row in data2]
x_data_2_peak2 = [float(row[0]) if is_float(row[0]) else np.nan for row in data2]
y_data_2_peak2 = [float(row[1]) if is_float(row[1]) else np.nan for row in data2]

x_data_3 = [float(row[0]) if is_float(row[0]) else np.nan for row in data3]
y_data_3 = [float(row[1]) if is_float(row[1]) else np.nan for row in data3]
x_data_3_peak1 = [float(row[0]) if is_float(row[0]) else np.nan for row in data3]
y_data_3_peak1 = [float(row[1]) if is_float(row[1]) else np.nan for row in data3]
x_data_3_peak2 = [float(row[0]) if is_float(row[0]) else np.nan for row in data3]
y_data_3_peak2 = [float(row[1]) if is_float(row[1]) else np.nan for row in data3]

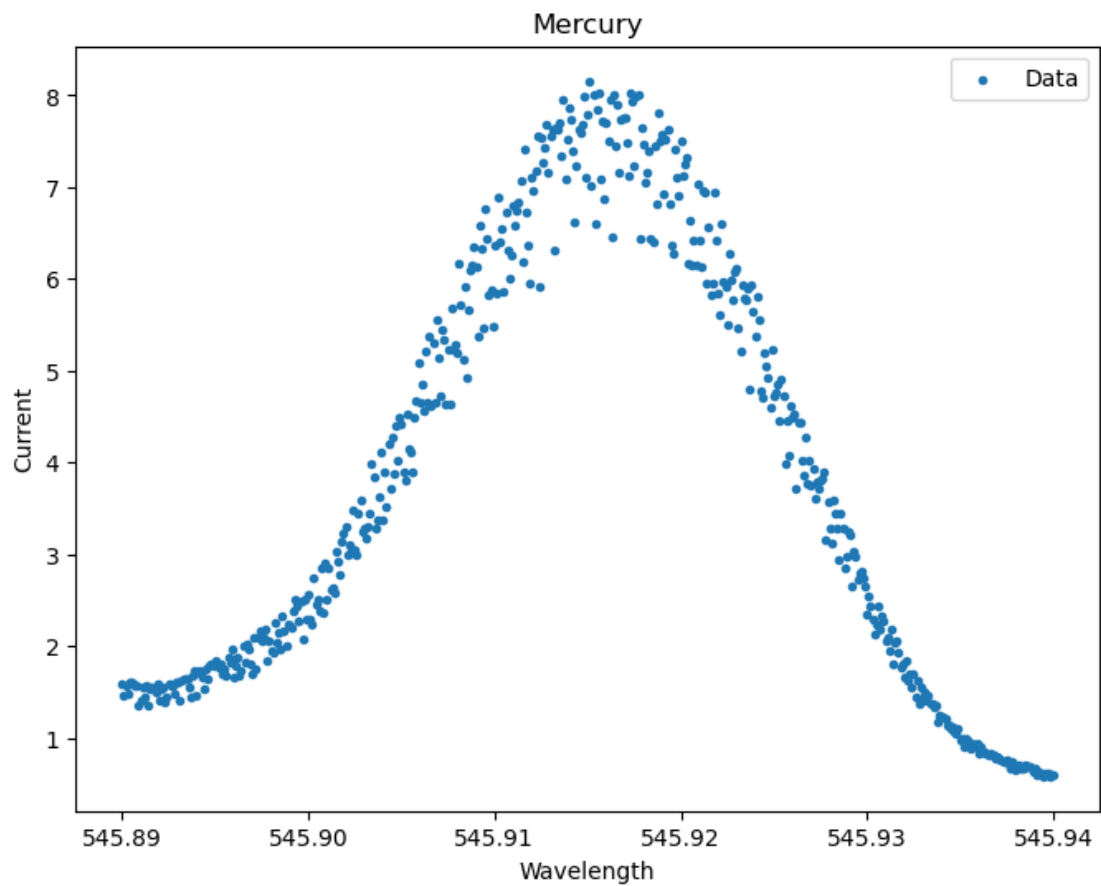
x_data_4 = [float(row[0]) if is_float(row[0]) else np.nan for row in data4]
y_data_4 = [float(row[1]) if is_float(row[1]) else np.nan for row in data4]
x_data_4_peak1 = [float(row[0]) if is_float(row[0]) else np.nan for row in data4]
y_data_4_peak1 = [float(row[1]) if is_float(row[1]) else np.nan for row in data4]
x_data_4_peak2 = [float(row[0]) if is_float(row[0]) else np.nan for row in data4]
y_data_4_peak2 = [float(row[1]) if is_float(row[1]) else np.nan for row in data4]

x_data_5 = [float(row[0]) if is_float(row[0]) else np.nan for row in data5]
y_data_5 = [float(row[1]) if is_float(row[1]) else np.nan for row in data5]
```

## Mercury Calibration

In [24]: `# plot all data`

```
plt.figure(figsize=(8, 6))
plt.scatter(x_data_m, y_data_m, label='Data', marker='.')
plt.xlabel('Wavelength')
plt.ylabel('Current')
plt.title('Mercury')
plt.legend()
plt.show()
```



```

In [25]: ▶ #x_min = 545.901
#x_max = 545.93
A1 = 7
B1 = 545.91
C1 = 0.01
D = 1

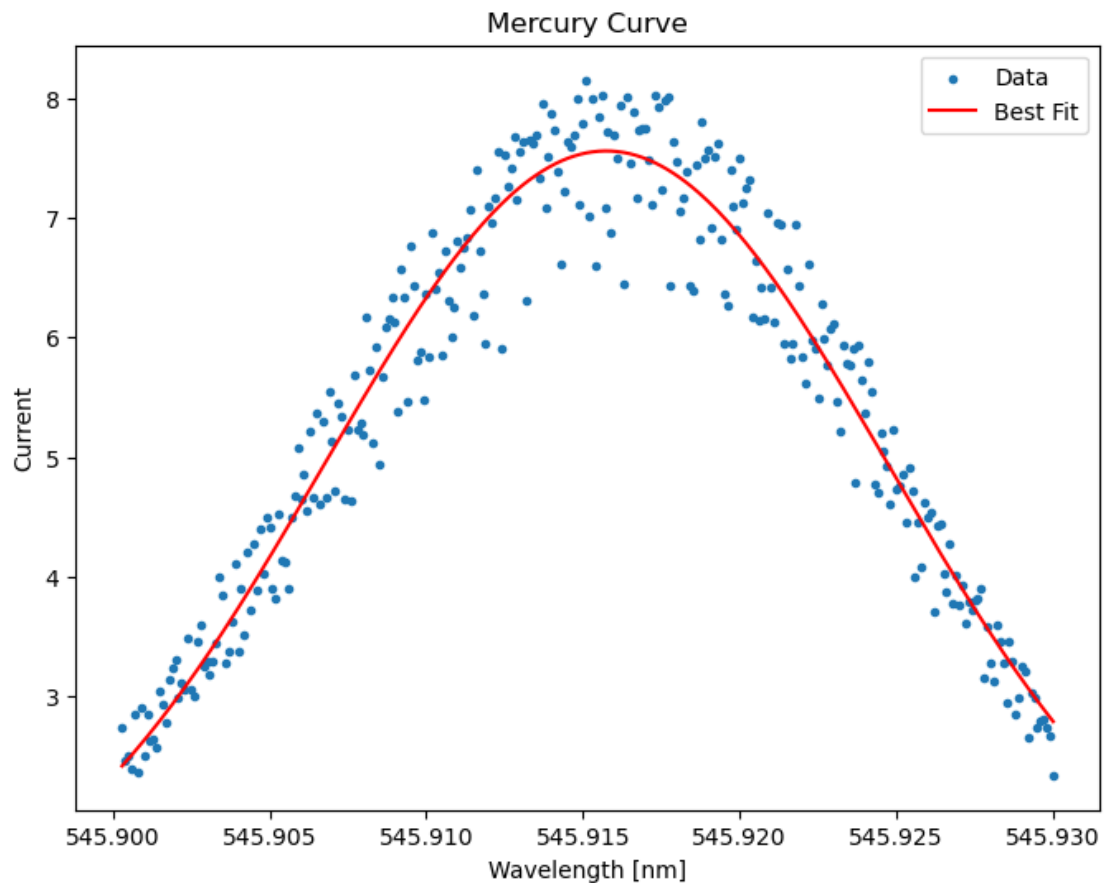
params, covariance = curve_fit(fgaussian, x_data_m_Peak1, y_data_m_Peak1,
                               p0=[A1, B1, C1, D])

A1_fit, B1_fit, C1_fit, D_fit = params
uncert = np.sqrt(np.diag(covariance))

plt.figure(figsize=(8, 6))
plt.scatter(x_data_m_Peak1, y_data_m_Peak1, label='Data', marker='.')
plt.plot(x_data_m_Peak1, fgaussian(x_data_m_Peak1, *params), label='Best Fit')
plt.xlabel('Wavelength [nm]')
plt.ylabel('Current')
plt.title('Mercury Curve')
plt.legend()
plt.show()

print('Peak 1 (0.081 MeV):')
print()
print(f'A1 = {A1_fit:.8f} ± {uncert[0]:.8f}')
print(f'B1 = {B1_fit:.8f} ± {uncert[1]:.8f}')
print(f'C1 = {C1_fit:.8f} ± {uncert[2]:.8f}')
print(f'D = {D_fit:.8f} ± {uncert[3]:.8f}')

```

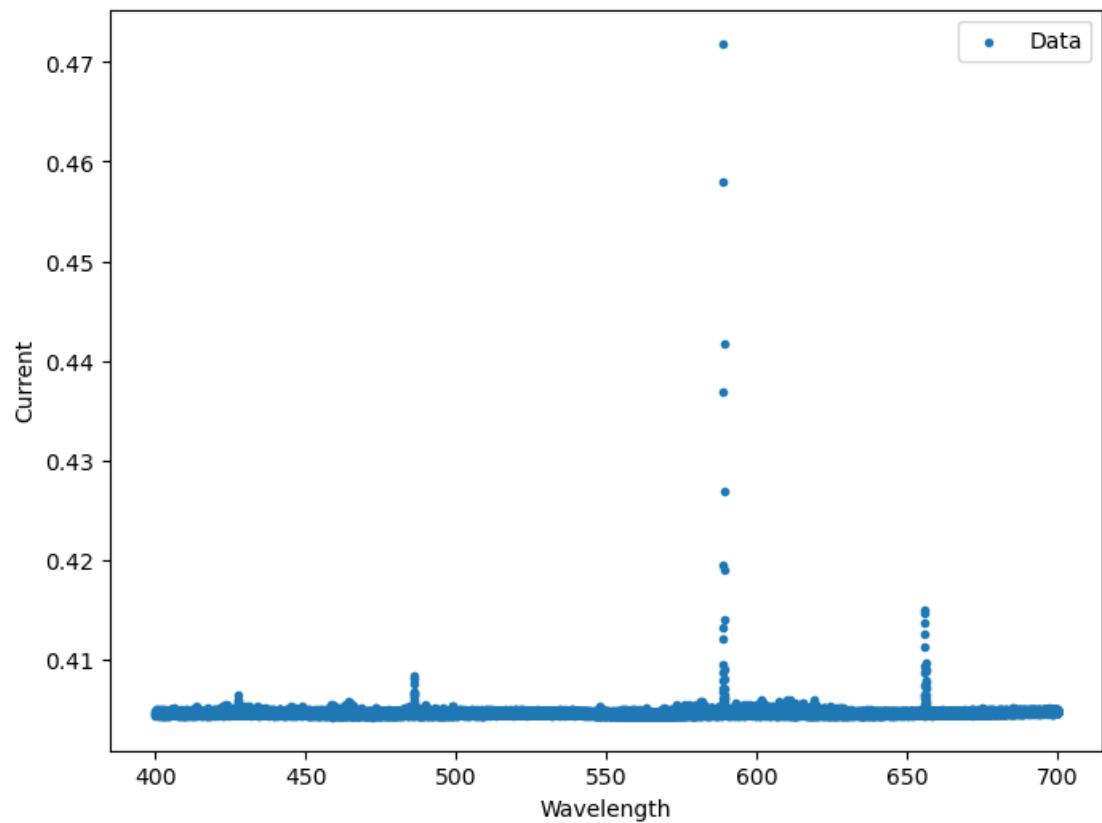


Peak 1 (0.081 MeV):

A1 =  $6.68835828 \pm 0.31518634$   
B1 =  $545.91574142 \pm 0.00006202$   
C1 =  $0.00900771 \pm 0.00038161$   
D =  $0.87393753 \pm 0.33408648$

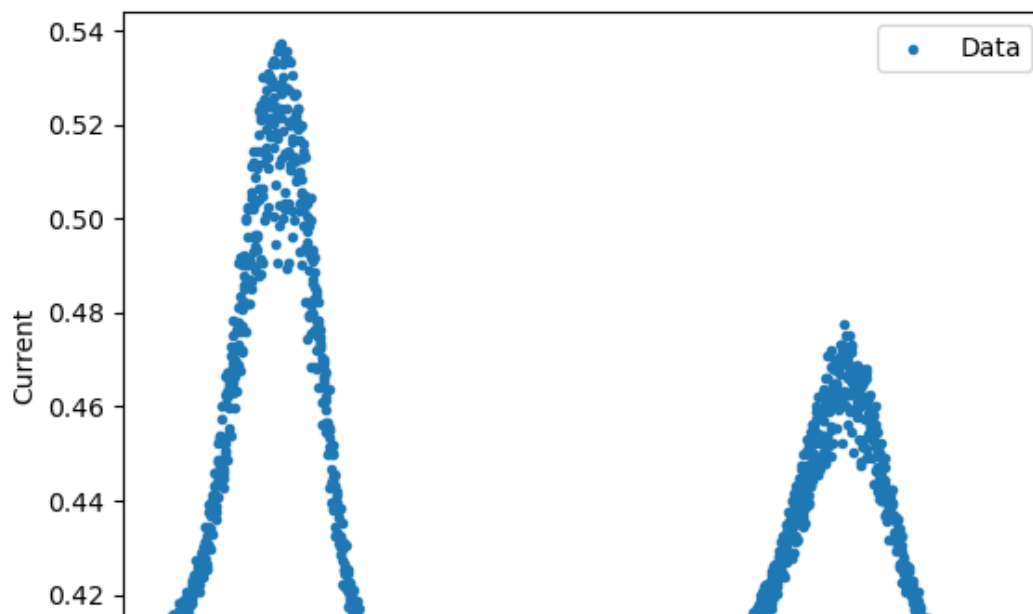
## Full Spectrum

```
In [26]: ▶ plt.figure(figsize=(8, 6))
plt.scatter(x_data_5, y_data_5, label='Data', marker='.')
plt.xlabel('Wavelength')
plt.ylabel('Current')
plt.title(' ')
plt.legend()
plt.show()
```



## 656.279 nm

```
In [27]: ▶ plt.scatter(x_data_1, y_data_1, label='Data', marker='.')  
plt.xlabel('Wavelength')  
plt.ylabel('Current')  
plt.title(' ')  
plt.legend()  
plt.show()
```



### **Peak 1**

(655.95, 656.05)

### **Peak 2**

(656.10, 656.25)



```

In [28]: ▶ A1 = 0.54
          B1 = 656
          C1 = .01
          D = 1

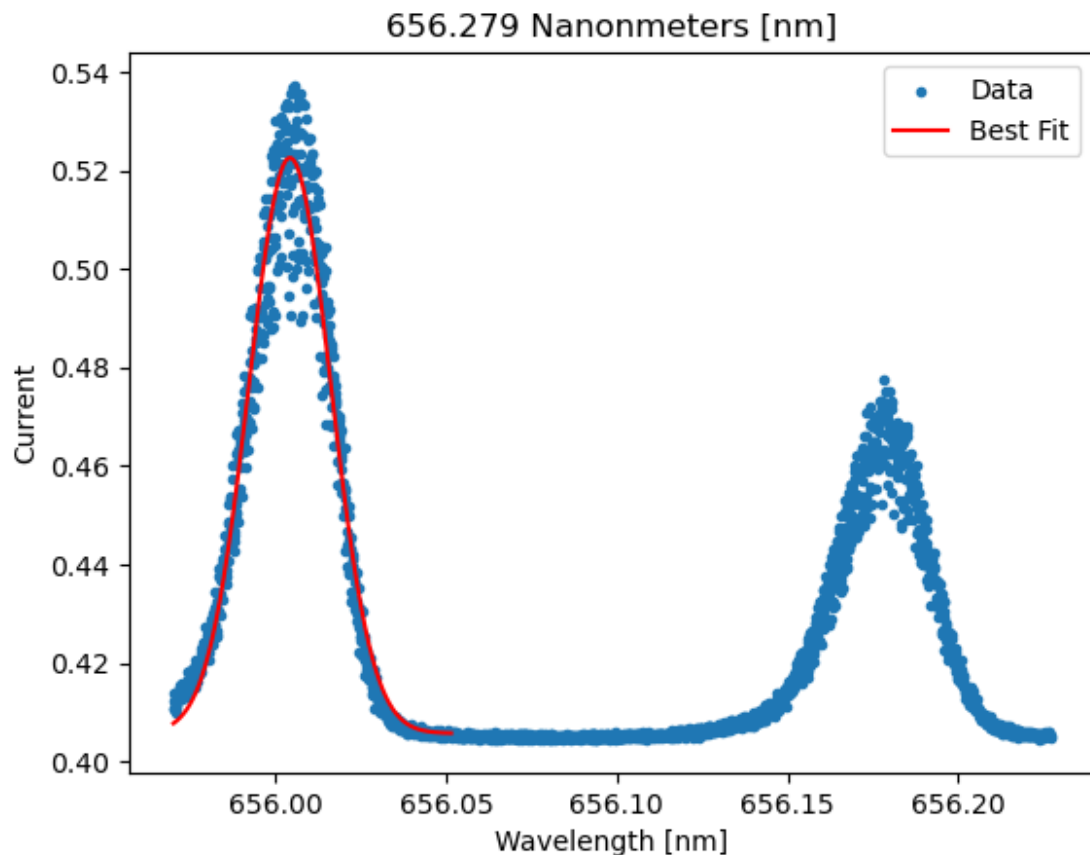
          params, covariance = curve_fit(fgaussian, x_data_1_peak1, y_data_1_peak1,
                                         p0=[A1, B1, C1, D])

          A1_fit, B1_fit, C1_fit, D_fit = params
          uncert = np.sqrt(np.diag(covariance))

          plt.scatter(x_data_1, y_data_1, label='Data', marker='.')
          plt.plot(x_data_1_peak1, fgaussian(x_data_1_peak1, *params), label='Best Fit')
          plt.xlabel('Wavelength [nm]')
          plt.ylabel('Current')
          plt.title('656.279 Nanometers [nm]')
          plt.legend()
          plt.show()

          print('656.279 Nanometers [nm]')
          print()
          print(f'A1 = {A1_fit:.8f} ± {uncert[0]:.8f}')
          print(f'B1 = {B1_fit:.8f} ± {uncert[1]:.8f}')
          print(f'C1 = {C1_fit:.8f} ± {uncert[2]:.8f}')

```



656.279 Nanometers [nm]

$$A1 = 0.11682109 \pm 0.00067805$$

$$B1 = 656.00413598 \pm 0.00006824$$

$$C1 = 0.01199412 \pm 0.00009666$$

```

In [29]: ▶ A1 = 0.54
          B1 = 656.156
          C1 = .05
          D = 1

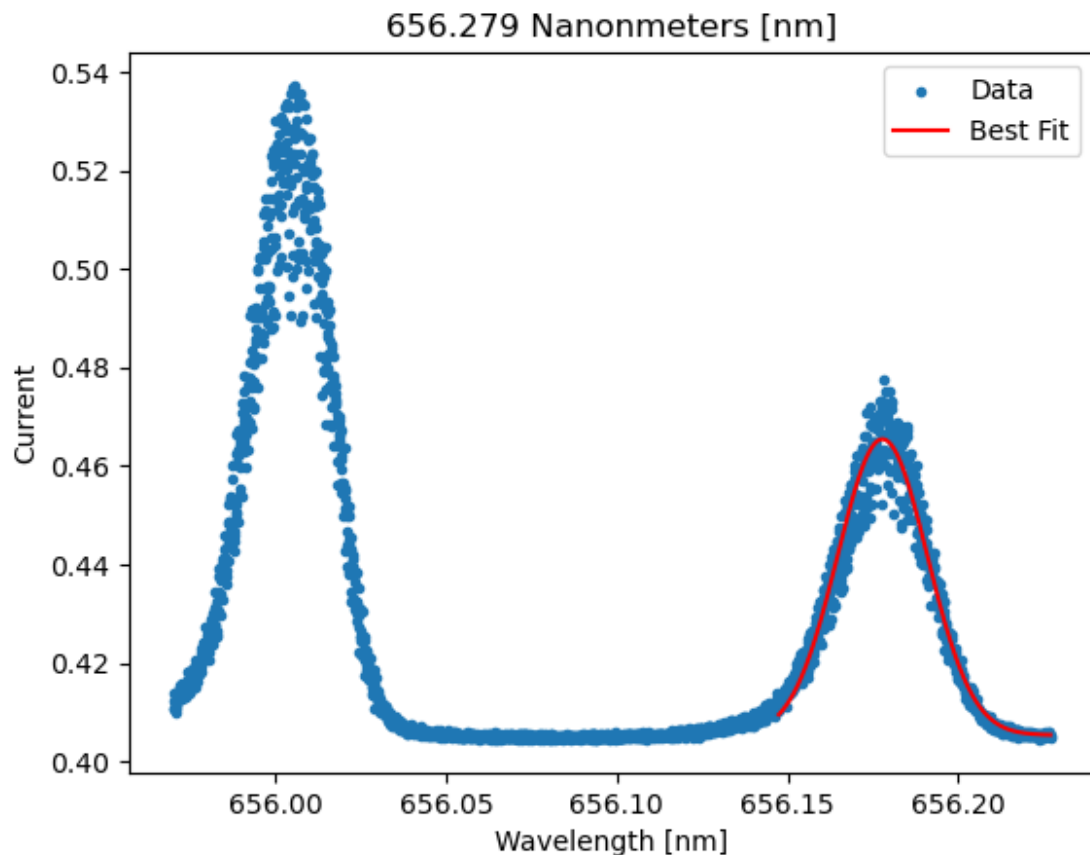
          params, covariance = curve_fit(fgaussian, x_data_1_peak2, y_data_1_peak2,
                                         p0=[A1, B1, C1, D])

          A1_fit, B1_fit, C1_fit, D_fit = params
          uncert = np.sqrt(np.diag(covariance))

          plt.scatter(x_data_1, y_data_1, label='Data', marker='.')
          plt.plot(x_data_1_peak2, fgaussian(x_data_1_peak2, *params), label='Best Fit')
          plt.xlabel('Wavelength [nm]')
          plt.ylabel('Current')
          plt.title('656.279 Nanometers [nm]')
          plt.legend()
          plt.show()

          print('656.279 Nanometers [nm]')
          print()
          print(f'A1 = {A1_fit:.8f} ± {uncert[0]:.8f}')
          print(f'B1 = {B1_fit:.8f} ± {uncert[1]:.8f}')
          print(f'C1 = {C1_fit:.8f} ± {uncert[2]:.8f}')

```



656.279 Nanometers [nm]

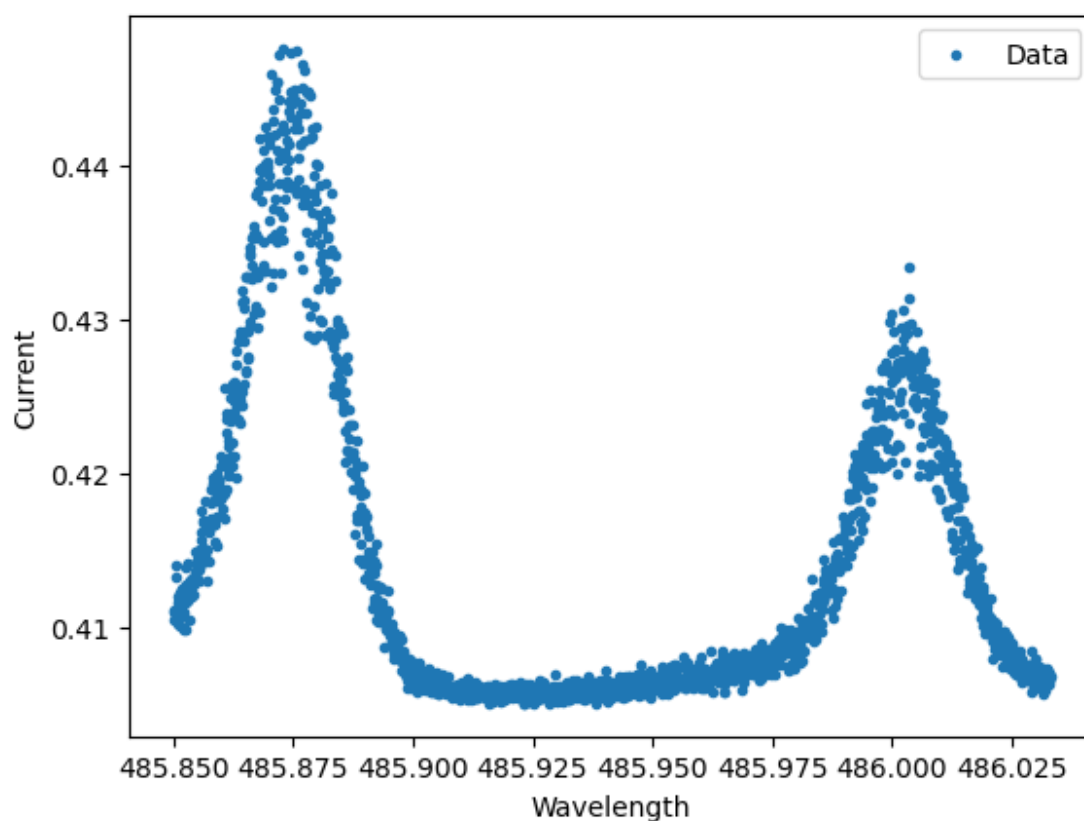
A1 =  $0.06007533 \pm 0.00037762$

B1 =  $656.17789794 \pm 0.00007609$

C1 =  $0.01316435 \pm 0.00011315$

## 486.135 nm

```
In [30]: ▶ plt.scatter(x_data_2, y_data_2, label='Data', marker='.')
plt.xlabel('Wavelength')
plt.ylabel('Current')
plt.title('')
plt.legend()
plt.show()
```



### Peak 1

(485, 485.9)

### Peak 2

(485.975, 486.025)

```

In [31]: ▶ A1 = 0.45
          B1 = 485.8
          C1 = .05
          D = 0

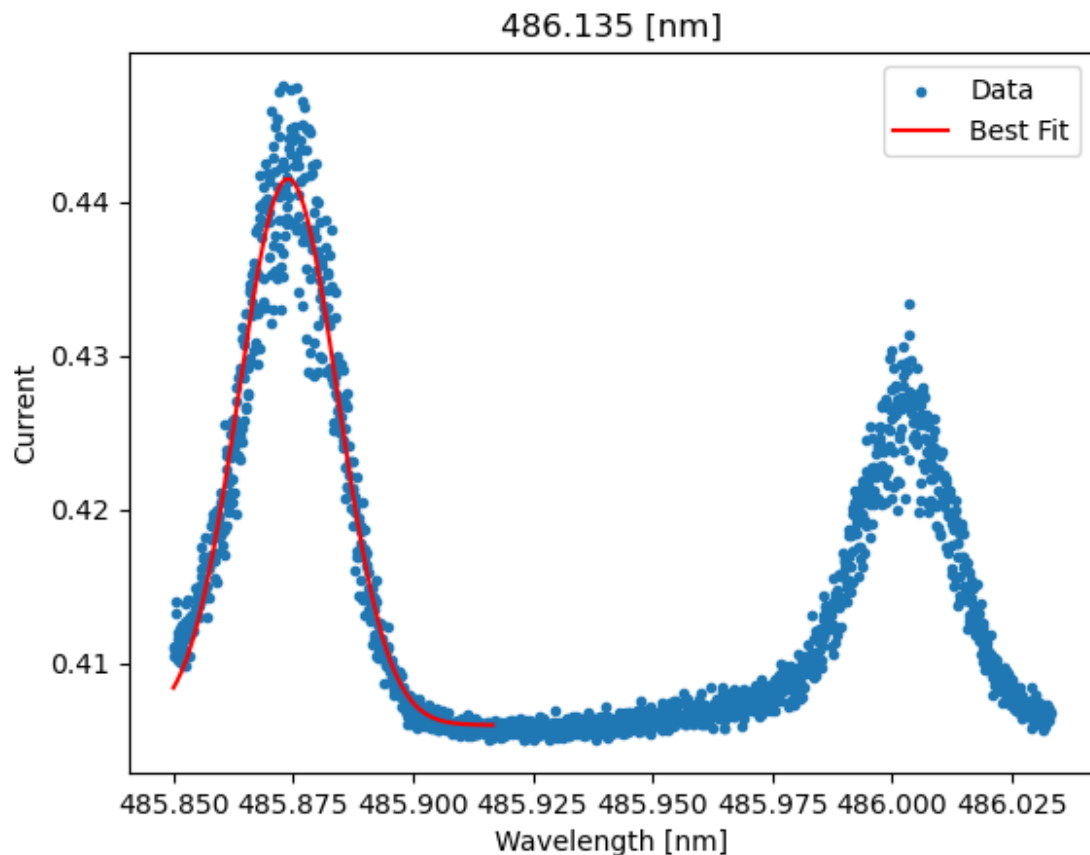
          params, covariance = curve_fit(fgaussian, x_data_2_peak1, y_data_2_peak1,
                                         p0=[A1, B1, C1, D])

          A1_fit, B1_fit, C1_fit, D_fit = params
          uncert = np.sqrt(np.diag(covariance))

          plt.scatter(x_data_2, y_data_2, label='Data', marker='.')
          plt.plot(x_data_2_peak1, fgaussian(x_data_2_peak1, *params), label='Best Fit')
          plt.xlabel('Wavelength [nm]')
          plt.ylabel('Current')
          plt.title('486.135 [nm]')
          plt.legend()
          plt.show()

          print('486.135 [nm]')
          print()
          print(f'A1 = {A1_fit:.8f} ± {uncert[0]:.8f}')
          print(f'B1 = {B1_fit:.8f} ± {uncert[1]:.8f}')
          print(f'C1 = {C1_fit:.8f} ± {uncert[2]:.8f}')

```



486.135 [nm]

A1 =  $0.03550120 \pm 0.00025833$

B1 =  $485.87392625 \pm 0.00007253$

C1 =  $-0.01032817 \pm 0.00010056$

```

In [32]: ▶ A1 = 0.43
          B1 = 486
          C1 = .02
          D = 0

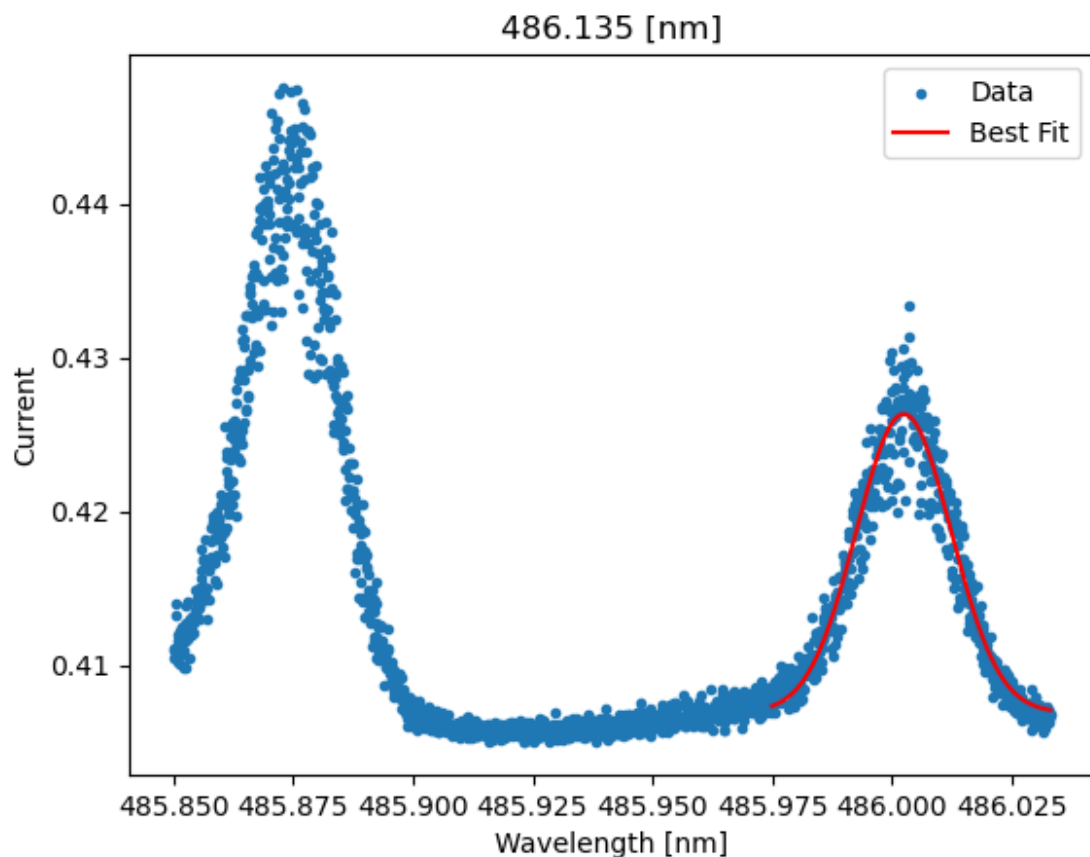
          params, covariance = curve_fit(fgaussian, x_data_2_peak2, y_data_2_peak2,
                                         p0=[A1, B1, C1, D])

          A1_fit, B1_fit, C1_fit, D_fit = params
          uncert = np.sqrt(np.diag(covariance))

          plt.scatter(x_data_2, y_data_2, label='Data', marker='.')
          plt.plot(x_data_2_peak2, fgaussian(x_data_2_peak2, *params), label='Best Fit')
          plt.xlabel('Wavelength [nm]')
          plt.ylabel('Current')
          plt.title('486.135 [nm]')
          plt.legend()
          plt.show()

          print('486.135 [nm]')
          print()
          print(f'A1 = {A1_fit:.8f} ± {uncert[0]:.8f}')
          print(f'B1 = {B1_fit:.8f} ± {uncert[1]:.8f}')
          print(f'C1 = {C1_fit:.8f} ± {uncert[2]:.8f}')

```



486.135 [nm]

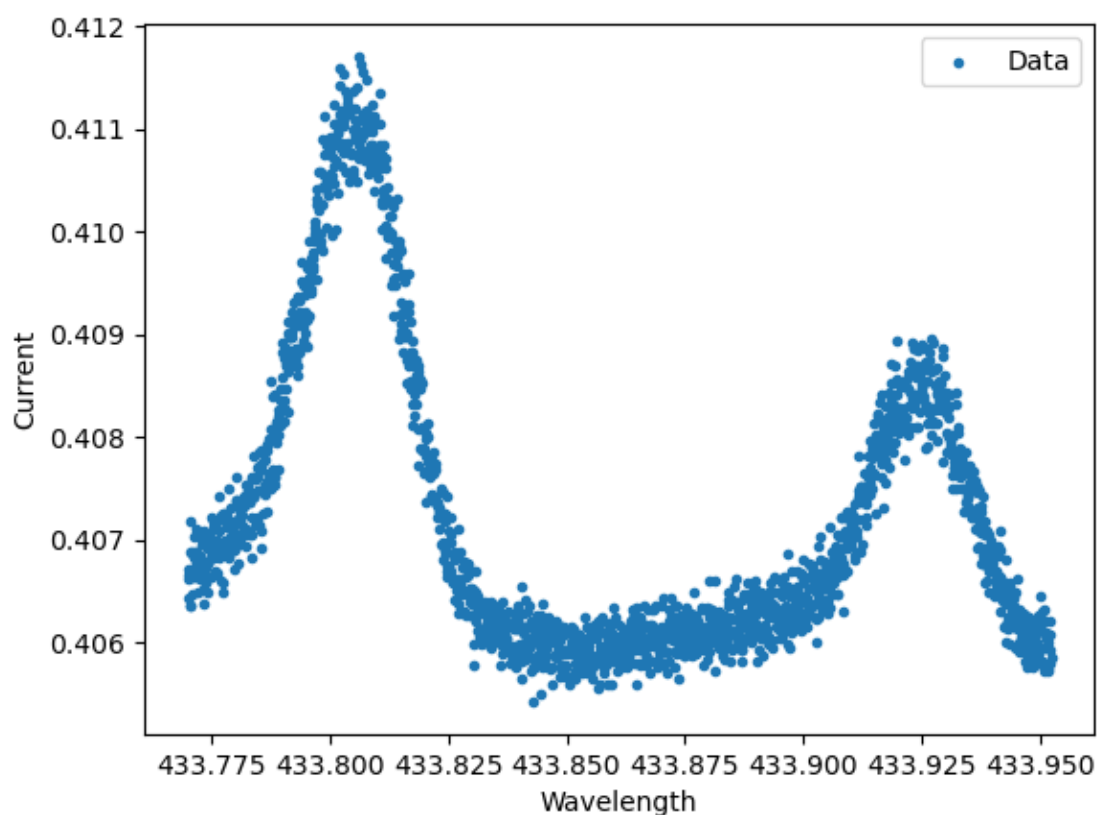
A1 =  $0.01939941 \pm 0.00020042$

B1 =  $486.00249349 \pm 0.00008774$

C1 =  $-0.00992384 \pm 0.00015021$

## 434.0472 nm

```
In [33]: ▶ plt.scatter(x_data_3, y_data_3, label='Data', marker='.')  
plt.xlabel('Wavelength')  
plt.ylabel('Current')  
plt.title('')  
plt.legend()  
plt.show()
```



### Peak 1

(433.775, 433.820)

### Peak 2

(433.900, 433.950)



```

In [34]: ▶ A1 = 0.412
          B1 = 433.8
          C1 = .1
          D = 0

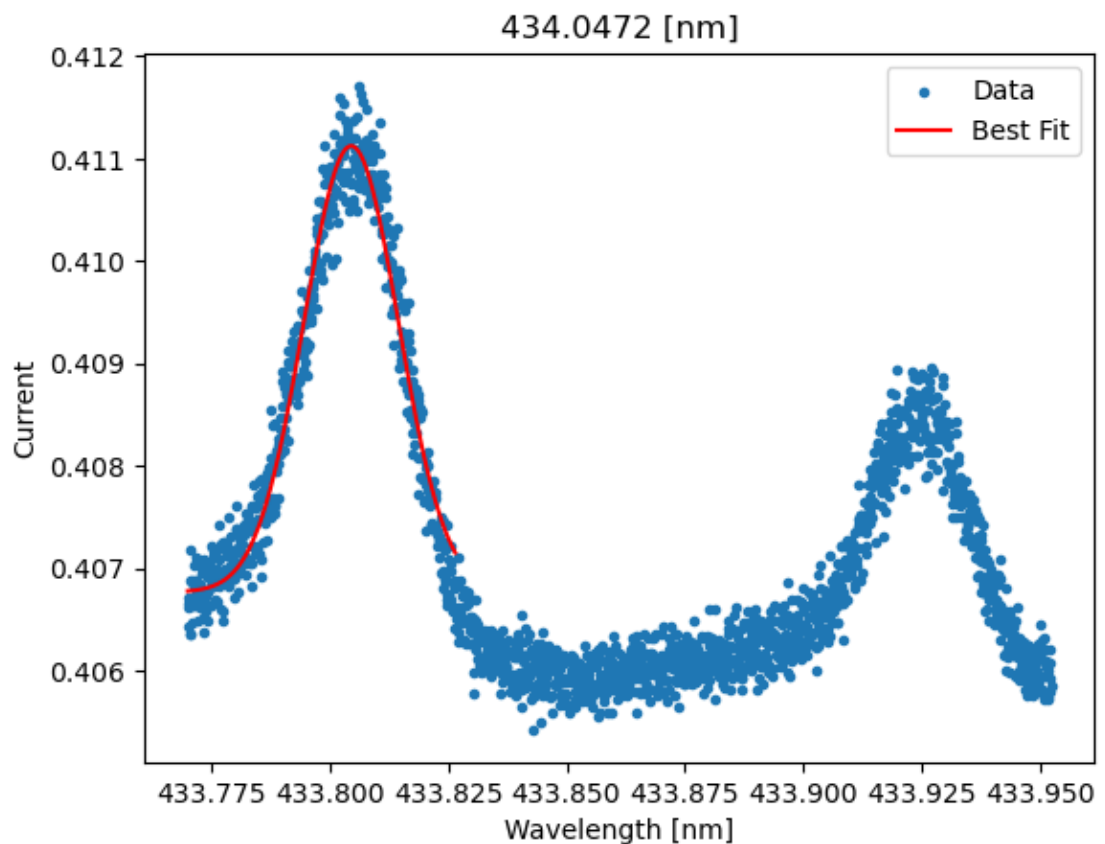
          params, covariance = curve_fit(fgaussian, x_data_3_peak1, y_data_3_peak1,
                                         p0=[A1, B1, C1, D])

          A1_fit, B1_fit, C1_fit, D_fit = params
          uncert = np.sqrt(np.diag(covariance))

          plt.scatter(x_data_3, y_data_3, label='Data', marker='.')
          plt.plot(x_data_3_peak1, fgaussian(x_data_3_peak1, *params), label='Best Fit')
          plt.xlabel('Wavelength [nm]')
          plt.ylabel('Current')
          plt.title('434.0472 [nm]')
          plt.legend()
          plt.show()

          print('434.0472 [nm]')
          print()
          print(f'A1 = {A1_fit:.8f} ± {uncert[0]:.8f}')
          print(f'B1 = {B1_fit:.8f} ± {uncert[1]:.8f}')
          print(f'C1 = {C1_fit:.8f} ± {uncert[2]:.8f}')

```



434.0472 [nm]

A1 = 0.00435986 ± 0.00003734

B1 = 433.80452053 ± 0.00007416

C1 = 0.00997645 ± 0.00011759

```

In [35]: ▶ A1 = 0.43
          B1 = 433.925
          C1 = .02
          D = 0

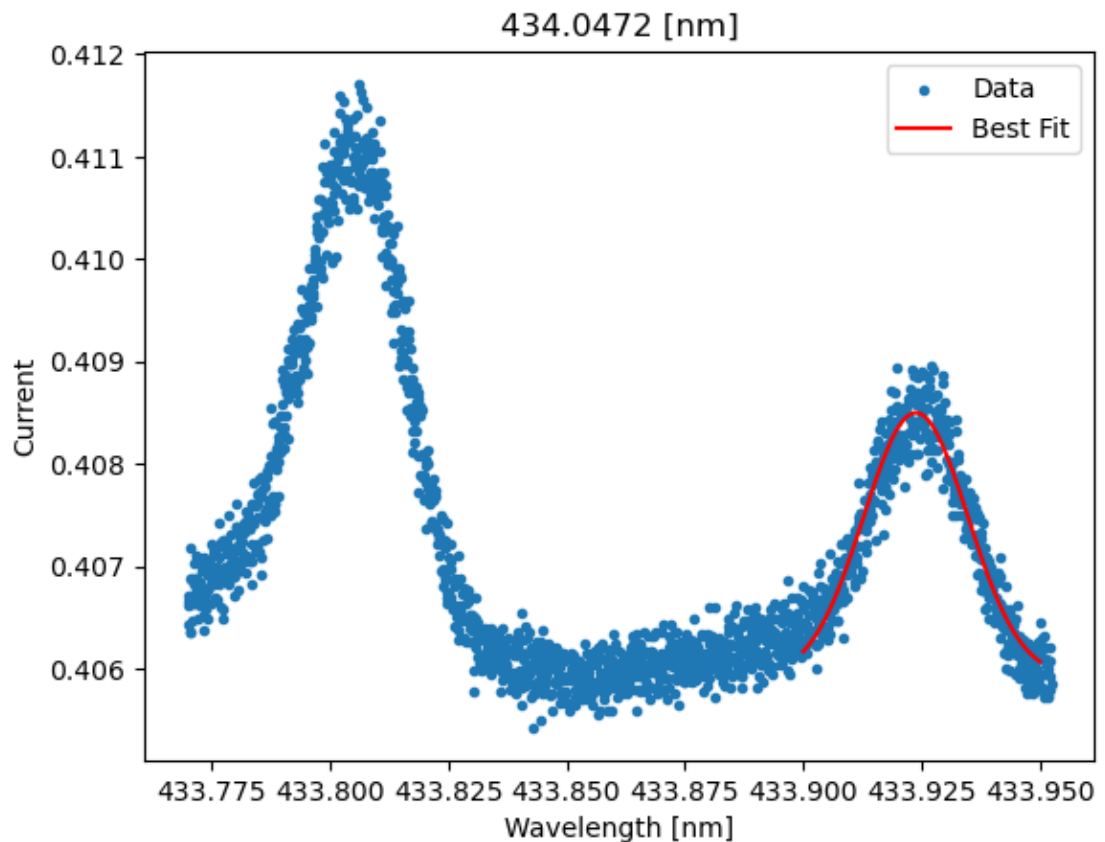
          params, covariance = curve_fit(fgaussian, x_data_3_peak2, y_data_3_peak2,
                                         p0=[A1, B1, C1, D])

          A1_fit, B1_fit, C1_fit, D_fit = params
          uncert = np.sqrt(np.diag(covariance))

          plt.scatter(x_data_3, y_data_3, label='Data', marker='.')
          plt.plot(x_data_3_peak2, fgaussian(x_data_3_peak2, *params), label='Best Fit')
          plt.xlabel('Wavelength [nm]')
          plt.ylabel('Current')
          plt.title('434.0472 [nm]')
          plt.legend()
          plt.show()

          print('434.0472 [nm]')
          print()
          print(f'A1 = {A1_fit:.8f} ± {uncert[0]:.8f}')
          print(f'B1 = {B1_fit:.8f} ± {uncert[1]:.8f}')
          print(f'C1 = {C1_fit:.8f} ± {uncert[2]:.8f}')

```



434.0472 [nm]

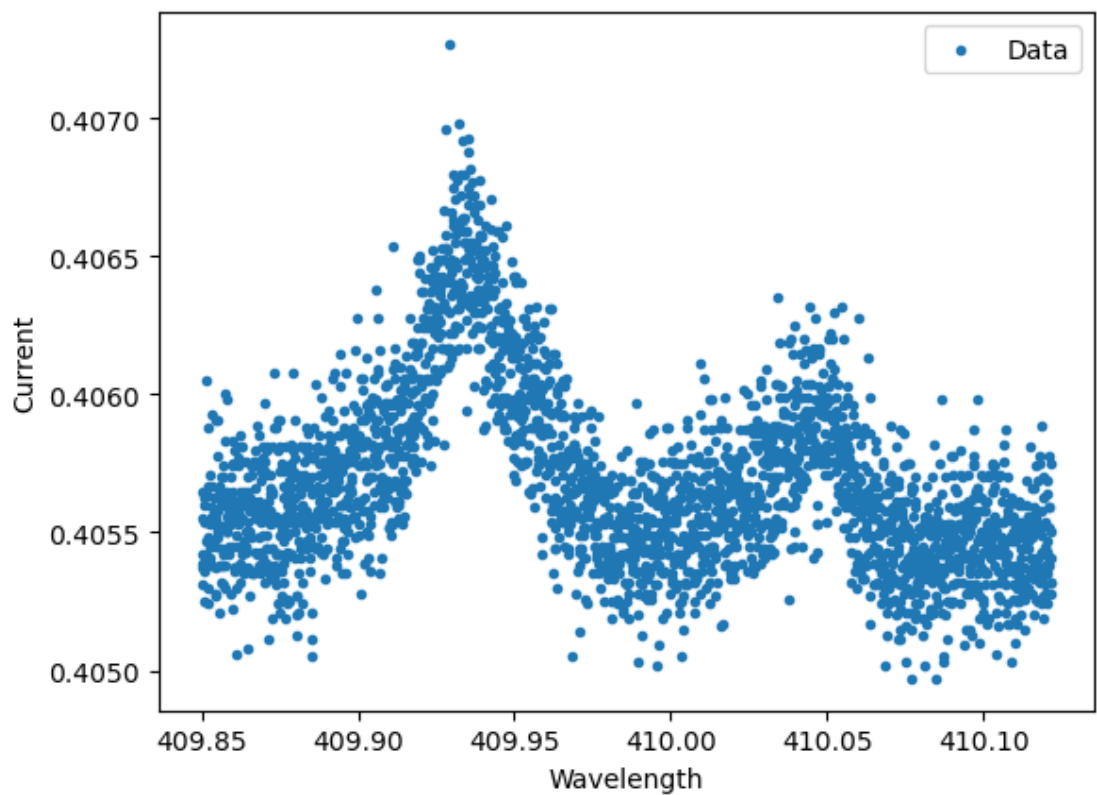
A1 =  $0.00258533 \pm 0.00005235$

B1 =  $433.92375671 \pm 0.00010745$

C1 =  $0.01102500 \pm 0.00029061$

## 410.1734 nm

```
In [36]: ▶ plt.scatter(x_data_4, y_data_4, label='Data', marker='.')  
plt.xlabel('Wavelength')  
plt.ylabel('Current')  
plt.title('')  
plt.legend()  
plt.show()
```



```

In [37]: ▶ A1 = 0.407
          B1 = 409.925
          C1 = .01
          D = 0

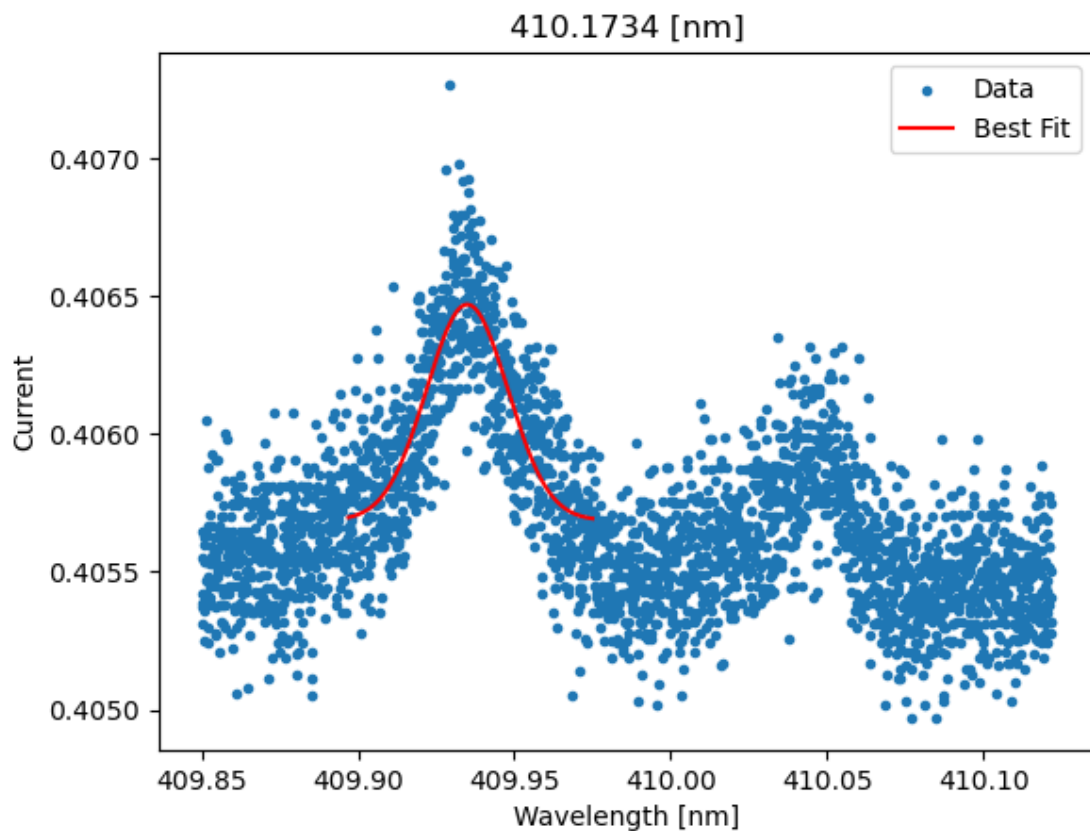
          params, covariance = curve_fit(fgaussian, x_data_4_peak1, y_data_4_peak1,
                                         p0=[A1, B1, C1, D])

          A1_fit, B1_fit, C1_fit, D_fit = params
          uncert = np.sqrt(np.diag(covariance))

          plt.scatter(x_data_4, y_data_4, label='Data', marker='.')
          plt.plot(x_data_4_peak1, fgaussian(x_data_4_peak1, *params), label='Best Fit')
          plt.xlabel('Wavelength [nm]')
          plt.ylabel('Current')
          plt.title('410.1734 [nm]')
          plt.legend()
          plt.show()

          print('410.1734 [nm]')
          print()
          print(f'A1 = {A1_fit:.8f} ± {uncert[0]:.8f}')
          print(f'B1 = {B1_fit:.8f} ± {uncert[1]:.8f}')
          print(f'C1 = {C1_fit:.8f} ± {uncert[2]:.8f}')

```



410.1734 [nm]

```

A1 = 0.00078410 ± 0.00002220
B1 = 409.93494744 ± 0.00032382
C1 = 0.01299440 ± 0.00053964

```

```

In [38]: ▶ A1 = 0.43
          B1 = 410.05
          C1 = .01
          D = 0

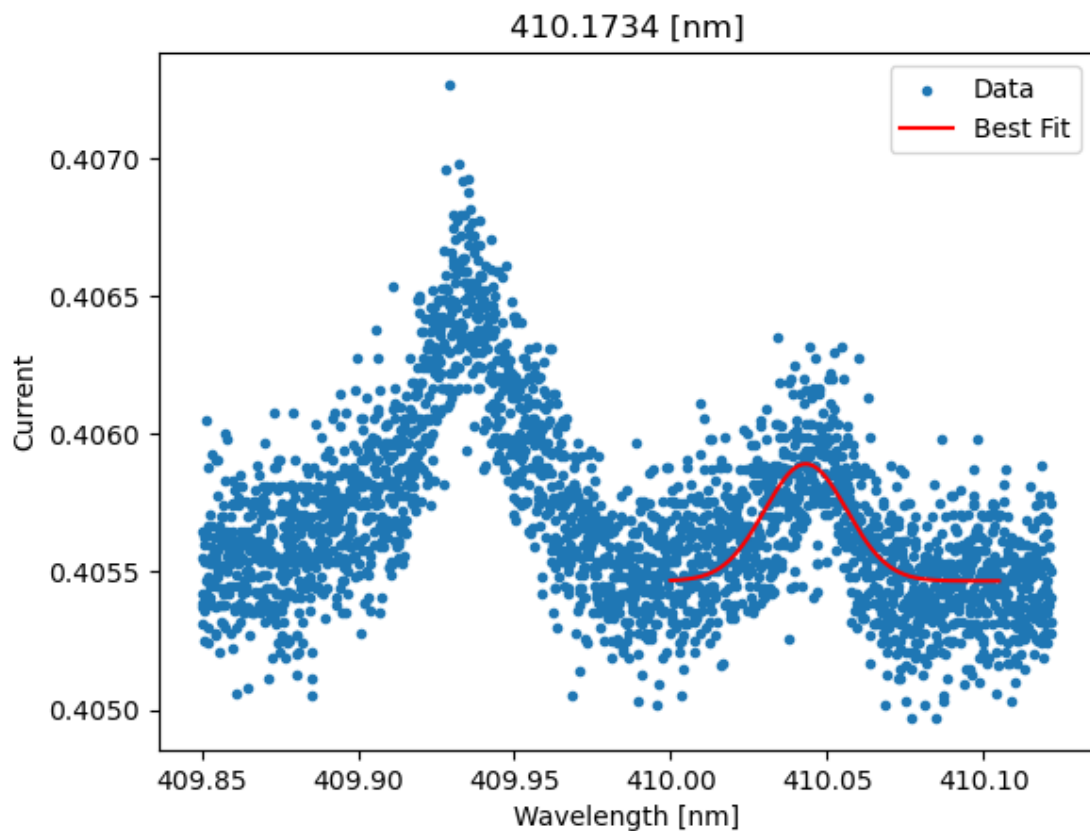
          params, covariance = curve_fit(fgaussian, x_data_4_peak2, y_data_4_peak2,
                                         p0=[A1, B1, C1, D])

          A1_fit, B1_fit, C1_fit, D_fit = params
          uncert = np.sqrt(np.diag(covariance))

          plt.scatter(x_data_4, y_data_4, label='Data', marker='.')
          plt.plot(x_data_4_peak2, fgaussian(x_data_4_peak2, *params), label='Best Fit')
          plt.xlabel('Wavelength [nm]')
          plt.ylabel('Current')
          plt.title('410.1734 [nm]')
          plt.legend()
          plt.show()

          print('410.1734 [nm]')
          print()
          print(f'A1 = {A1_fit:.8f} ± {uncert[0]:.8f}')
          print(f'B1 = {B1_fit:.8f} ± {uncert[1]:.8f}')
          print(f'C1 = {C1_fit:.8f} ± {uncert[2]:.8f}')

```



410.1734 [nm]

A1 = 0.00042482 ± 0.00001670  
 B1 = 410.04316769 ± 0.00053846  
 C1 = 0.01310418 ± 0.00068792

## Plot

$$\frac{1}{\lambda} = R \left( \frac{1}{2^2} - \frac{1}{n_i^2} \right)$$

$$y = \frac{1}{\lambda} \quad x = \frac{1}{n_i^2}$$

$$y = R \left( \frac{1}{2^2} - x \right) \rightarrow y = \frac{R}{4} - Rx$$

Note: Rydberg Constant for an Infinitely Massive Nucleus

$$R_{\infty} = \frac{1}{hc} \frac{E_0}{2} = \frac{1}{2} \frac{1}{hc} a^2 m_e c^2 = \frac{1}{1240 \text{ eV nm}} \frac{1}{137^2} (0.511 \times 10^6 \text{ eV}) = 0.01097 \text{ nm}^{-1}$$

Rydberg Constant for Deuterium and Hydrogen:

$$R_D = \frac{\mu_D}{m_e} R_{\infty}$$

$$R_H = \frac{\mu_H}{m_e} R_{\infty}$$

You can use these to find the reduced mass/electron mass ratio by dividing the specific Rydberg constant by the Rydberg constant for an infinitely massive nucleus.

$$\text{e.g.} \quad R_D = \frac{\mu_D}{m_e} R_{\infty}$$

$$\frac{\mu_D}{m_e} = \frac{R_D}{R_{\infty}}$$

```

In [39]: ▶ R_inf = 0.01097 # nm^-1

i = np.array([1.000224458, 1.000186905,
              1.000228641, 1.000228751]) # indices of refraction

ld = np.array([656.00413598, 485.87392625,
              433.80452053, 409.93494744]) # wavelengths for deuterium

yd = 1/(ld*i) # y for deuterium

ud = np.array([0.00006824, 0.00007253,
              0.00007416, 0.00032382]) # deuterium uncert

lh = np.array([656.17789794, 486.00249349,
              433.92375671, 410.04316769]) # wavelengths for hydrogen

yh = 1/(lh*i) # y for hydrogen

uh = np.array([0.00007609, 0.00008774,
              0.00010745, 0.00053846]) # hydrogen uncert

n = np.array([3, 4, 5, 6]) # energy level number

x_n = 1/n**2 # x

yd_u = 1/(i*(ld-ud)) # upper end of the error for deuterium

yh_u = 1/(i*(lh-uh)) # upper end of error for hydrogen

yd_err = abs(yd - yd_u) # error for y deuterium
yh_err = abs(yh - yh_u) # error for y hydrogen

print(yd)
print()
print(yd_u)
print()
print(yd_err)
print()
print(yh_err)

[0.00152404 0.00205776 0.00230466 0.00243885]

[0.00152404 0.00205776 0.00230466 0.00243886]

[1.58536198e-10 3.07177492e-10 3.93987427e-10 1.92652562e-09]

[1.76679828e-10 3.71397990e-10 5.70532496e-10 3.20180975e-09]

```

## Deuterium



```

In [40]: slope, intercept, r_value, p_value, std_err = stats.linregress(x_n, yd)

result = stats.linregress(x_n, yd)

data_labels = ['n = 3', 'n = 4',
               'n = 5', 'n = 6']

for x, y, label in zip(x_n, yd, data_labels):
    plt.text(x, y, label, fontsize=9, ha='left', va='top')

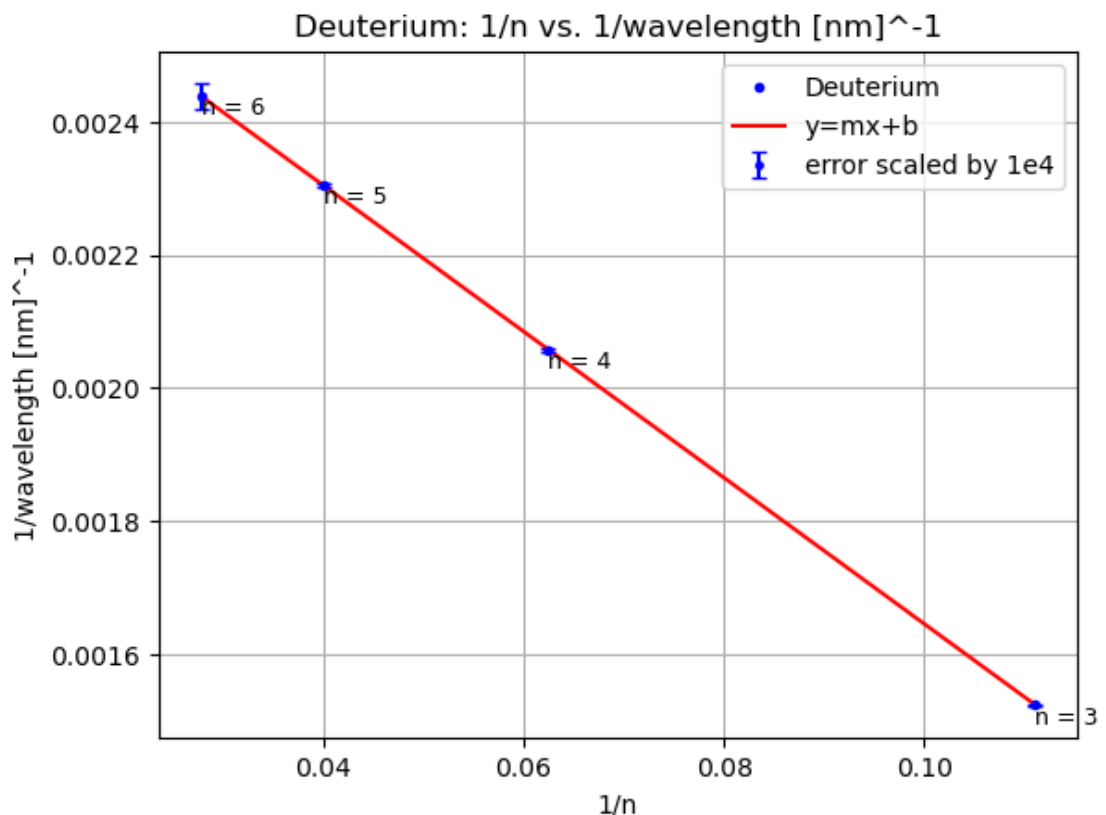
res = stats.linregress(x_n, yd)
plt.plot(x_n, yd, '.', color='b', label='Deuterium')
plt.plot(x_n, res.intercept + res.slope*x_n, 'r', label='y=mx+b')
plt.errorbar(x_n, yd, yerr=1e4*yd_err, fmt='.', color='b',
             label='error scaled by 1e4', markersize=5, capsize=3)

plt.xlabel('1/n')
plt.ylabel('1/wavelength [nm]^(-1)')
plt.title('Deuterium: 1/n vs. 1/wavelength [nm]^(-1)')
plt.grid(True)
plt.legend()
plt.show()

# chi_sq = np.sum((yd-(res.intercept+res.slope*x_n))**2/yd_err**2)

print(f"Linear fit: y = ({res.slope:.8f})+/-({result.stderr:.8f})x + ({res.intercept:.8f})")
#print()
#print(f'Chi-squared: {chi_sq}')

```



Linear fit:  $y = (-0.01097763 \pm 0.00000087)x + (0.00274380 \pm 0.00000006)$

```
In [41]: ▶ Rd = 0.01097763 # nm-1
sigma_Rd = 0.00000087
Rd_r = 0.999728

print(f'Rydberg constant for Deuterium: {Rd}')
print()
print(f'Calculated reduced mass/electron mass ratio: {Rd/R_inf}')
print()
print(f'Actual reduced mass/electron mass ratio: {Rd_r}')
print()
print(f'Absolute error for reduced mass/electron mass ratio: {abs(Rd - Rd_r)')
```

Rydberg constant for Deuterium: 0.01097763

Calculated reduced mass/electron mass ratio: 1.0006955332725616

Actual reduced mass/electron mass ratio: 0.999728

Absolute error for reduced mass/electron mass ratio: 0.98875037

## Hydrogen

```

In [42]: slope, intercept, r_value, p_value, std_err = stats.linregress(x_n, yh)

result = stats.linregress(x_n, yh)

data_labels = ['n = 3', 'n = 4',
               'n = 5', 'n = 6']

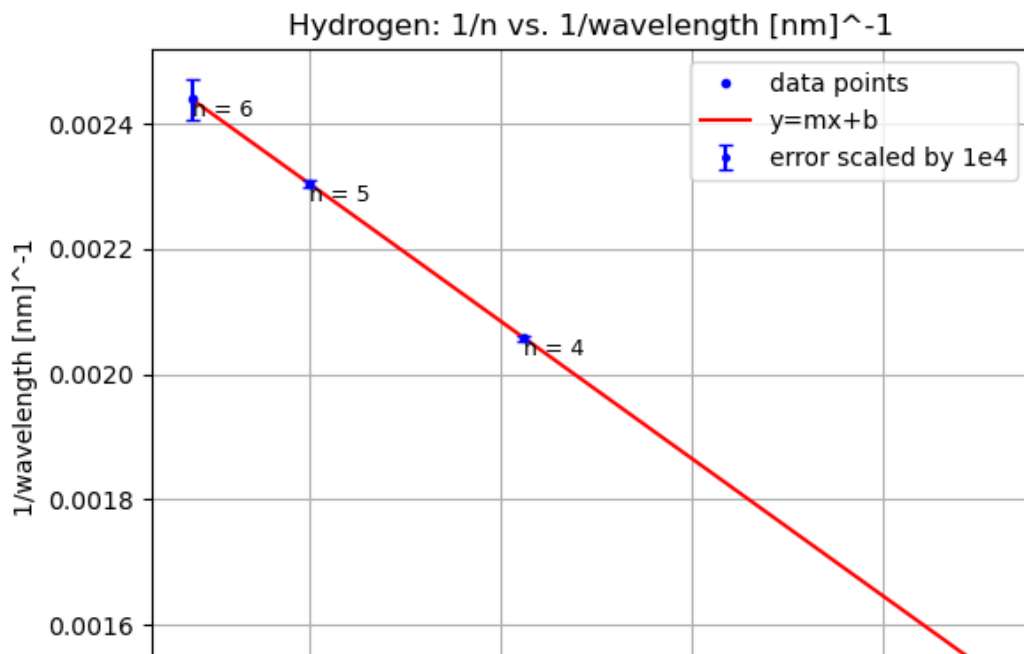
for x, y, label in zip(x_n, yh, data_labels):
    plt.text(x, y, label, fontsize=9, ha='left', va='top')

res = stats.linregress(x_n, yh)
plt.plot(x_n, yh, '.', color='blue', label='data points')
plt.plot(x_n, res.intercept + res.slope*x_n, 'r', label='y=mx+b')
plt.errorbar(x_n, yh, yerr=1e4*yh_err, fmt='.', color='blue',
             label='error scaled by 1e4', markersize=5, capsize=3)

plt.xlabel('1/n')
plt.ylabel('1/wavelength [nm]^(-1)')
plt.title('Hydrogen: 1/n vs. 1/wavelength [nm]^(-1)')
plt.grid(True)
plt.legend()
plt.show()

print(f"Linear fit: y = ({res.slope:.8f})+/-({result.stderr:.8f})x + ({res.intercept:.8f})")

```



```
In [43]: ▶ Rh = 0.01097462 # nm^-1
sigma_Rh = 0.00000100
Rh_r = 0.999457

print(f'Rydberg constant for Hydrogen: {Rh}')
print()
print(f'Reduced mass/electron mass ratio: {Rh/R_inf}')
print()
print(f'Actual reduced mass/electron mass ratio: {0.999457}')
print()
print(f'Absolute error for reduced mass/electron mass ratio: {abs(Rh - Rh_r)')
```

Rydberg constant for Hydrogen: 0.01097462

Reduced mass/electron mass ratio: 1.0004211485870556

Actual reduced mass/electron mass ratio: 0.999457

Absolute error for reduced mass/electron mass ratio: 0.98848238

$$\frac{R_D}{R_H} = \frac{\frac{\mu_D}{m_e} R_\infty}{\frac{\mu_H}{m_e} R_\infty} = \frac{\frac{1}{1 + \frac{m_e}{M_D}} R_\infty}{\frac{1}{1 + \frac{m_e}{M_H}} R_\infty} = \frac{1 + \frac{m_e}{M_H}}{1 + \frac{m_e}{M_D}} \approx \left(1 + \frac{m_e}{M_H}\right) \left(1 - \frac{m_e}{M_D}\right)$$

Neglecting second order terms.

$$\frac{R_D}{R_H} \approx 1 + m_e \left( \frac{1}{M_H} - \frac{1}{M_D} \right) = 1 + \frac{m_e}{M_H} \left( 1 - \frac{M_H}{M_D} \right)$$

$$\frac{m_e}{M_H} \approx \frac{\frac{R_D}{R_H} - 1}{1 - \frac{M_H}{M_D}}$$

From pre-lab writeup:

$$\frac{M_D}{M_H} = 1.999$$

$$\frac{m_e}{M_H} \approx \frac{\frac{R_D}{R_H} - 1}{1 - 1.999} = 2.001 \left( \frac{R_D}{R_H} - 1 \right)$$

```
In [60]: ▶ print(f'Ratio of the Rydberg constant for Deuterium and the Rydberg constant for Hydrogen: Rd/Rh = {Rd/Rh}')
print()
print(f'Electron/proton (Hydrogen nucleus) mass ratio: {2.001*((Rd/Rh)-1)}')
print()
print(f'Proton/electron mass ratio: {1/(2.001*((Rd/Rh)-1))}')
print()
print(f'Our proton/electron mass ratio is within {(abs(1/(2.001*((Rd/Rh)-1))-1836))}% the published value of 1836.'
```

Ratio of the Rydberg constant for Deuterium and the Rydberg constant for Hydrogen: Rd/Rh = 1.0002742691774296

Electron/proton (Hydrogen nucleus) mass ratio: 0.000548812624036559

Proton/electron mass ratio: 1822.1155203116928

Our proton/electron mass ratio is within 0.8% the published value of 1836.

In [ ]: ▶