Introduction to Data Analysis and Plotting with Jupyter

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The goal of this exercise is to familiarize the student with basic data manipulation and plotting using Jupyter notebooks. Many of the concepts are general and can be translated to other software applications. By the end of this exercise, you should be able to

- perform basic manipulations on data columns (formatting, transformations)
- perform simple mathematical operations over a set of data
- plot columns of data in a clear and visually pleasing manner
- perform simple statistical operations on a set of data
- · get a line-of-best-fit for multiple datasets
- import data from plain text files (CSV, TSV, DAT, etc.), like those created by many spectrometers and logging instruments

Introduction

Why do we plot data? Edward Tufte may have said it best:

At their best, graphics are instruments for reasoning about quantitative information. Often the most effective way to describe, explore, and summarize a set of numbers—even a very large set—is to look at pictures of those numbers.

A Little Bit About Plots

Plots (also called graphs or charts, though these are not *quite* the same thing) come in a variety of styles. The can be used to explore data as your are performing analyses or to convey results to a reader as part of a report. There are a variety of "standard" types which should cover your needs in this course. The main types we will use are:

Scatter plot Used to show correlation between two variables, each point generally represents an **independent** measurement.

Lines are **not** used to connect points, lines are instead used to represent a mathematical model fitting the data.

Line graph Indicates continuity in the *x* dimension, often suggests a single collection session for all data on the "line" (e.g., a UV-Vis or NMR spectrum). Individual points are generally not shown, though the data generally consists of individual collection points. Lines

should never be smoothed (or "splined"), as this treatment no longer accurately represents the data.

Really, a line graph is just a special case of a scatter plot where the independent variable is continuous (like a fluorescence or NMR spectrum).

Histogram Represents the distribution of data in a series of "bins", shows the frequency of a measured event. Closely related to density plots, in the same way scatter plots are related to line plots.

Here are the important parts of a plot, along with a brief description and guidelines for formatting.

Title Should be descriptive and represent all data in the plot.

Axes Should be clearly labeled (category, units, tick numbers). Should have an appropriate range (from slightly less than your minimum value to slightly more than your maximum value).

Fit line Should cover the whole spread of the data, but should not predict beyond the data range (you don't know anything about measurements in that region).

Legend Should clearly show which set of data is which, trend lines should clearly correlate to a given data set.

Basic Plotting and Linear Fits

Your first task is to take the data in the file data.txt and import it into a new Jupyter notebook.¹

As the first step, please use this link to clone the files into your Jupyter directory:

https://bit.ly/chem357-plot-intro

This link will direct your browser to the class JupyterHub and download a folder containing a number of resources you'll need for this exercise.

The folder raw_data contains a few datasets in various formats. The file extension doesn't always tell you what type of information is in a file and can sometimes lead you astray. For this exercise, the file data.txt contains a set of "tab-delimited" data, meaning that the values in a row are separated by tab characters. Another common format is "comma-delimited" or "comma-separated" data, such as the dataset in anscombe.csv. You will find that different instruments have different defaults for exporting data, but these two are the most common. Sometimes, a .dat extension might be used for a plaintext file to indicate 'data' without indicating the underlying format or delimiter type.

¹ Note: prior to performing the following exercises, you should run through the introductory documentation to Jupyter to familiarize yourself with the concepts and interface.

Make sure that you are located in the top folder for this lesson (i.e., inside of "357 plot intro") and open a new Jupyter template by

- 1. clicking on the "New Launcher" button at the top of the file browser panel, or
- 2. Selecting "File > New Launcher",

then clicking on the Template icon, select "pchem_templates" in the top box and "pchem report.ipynb" in the second box. You can also create a file from a new template directly by clicking on "File > New > Template". These methods are all equivalent, pick whichever you're most comfortable using. This will open a new Jupyter notebook with a series of cells pre-populated with information to get you started. Give this notebook a title (as instructed in the first cell), then click on "File > Rename Notebook..." and give the notebook a title following the instructions in the first cell.

In the notebook, execute the code cell under the "Library Import" section. "Libraries" are just collections of useful Python code that individuals or groups have created. They generally cover a specific area or hold to a theme. In our notebook template, we use the following:

NumPy A general purpose numerical library that speeds up number based operations, allows functions to operate on lists of numbers, and gives a large collection of useful mathematical functions (things like polynomial fitting, matrix operations, and common algebraic and trigonometric functions). The core functionality is based around NumPy arrays, very fast n-dimensional arrays of data (a core requirement for scientific computation).

SciPy A fundamental library for scientific computing, builds on the NumPy library but includes numeric integration routines, non-linear curve fitting routines, interpolation, numerical optimization, and statistics.

Pandas A data management library that offers functions for importing, tidying, organizing, and filtering data using the DataFrame construct. Excellent resource for 2-dimensional data, starts breaking down with more dimensions (look at the xarray package, which is built on top of Pandas, for this purpose).

Matplotlib A comprehensive library for creating static, animated, and interactive visualizations in Python. It has become the de-facto standard library for plot visualizations in Python.

Seaborn A collection of routines and formats built on top of Matplotlib that provides a high-level interface for drawing attractive and informative statistical graphics.

Now, create a new cell under the Data import section. Import the data from data.txt by using the import function of the "Pandas" (pd) library:

```
data = pd.read csv("raw data/data.txt", sep="\t")
```

This will take the contents of data.txt and turn it into an object called a "DataFrame" with the name data. The sep="\t" argument is required because our data is "tab-separated", rather than "commaseparated", which is the default (as suggested by the function name read csv). When in doubt, specify the field separator when you import data with python and the pandas library.

To view your new object, just type data on a new line and press Shift + Enter to evaluate the cell. By default code cells will display the results of the final line of the cell (if there is anything to display).³ You should end up with two columns of data with 21 data points in each column. Once you have the data columns, create a new scatter plot from the data. While the Pandas library automatically incorporates basic plotting functions from the Matplotlib library, customizing the plot can often be difficult.⁴ While you are exploring data, the default appearance is fine, but you'll probably want to start customizing the appearance of plots when you get them ready for publication.

Adding data to the plot function is done in the following manner:

```
plt.plot(x1, y1, ['fmt1'], x2, y2, ['fmt2'], ...,

    data=data-source)
```

This method is useful if you know the structure of the data ahead of time (and when you have a clear index column, such as a time index, wavelength, or frequency). The second way to fix the plot is to define the x and y data in the arguments of the plot() function. The plot() function takes a number of arguments (you can see them quickly by pressing | Shift | + | Tab | inside of the parentheses), including arguments for features like plot type (e.g., kind="scatter"), plot title (e.g., title="My Plot"), and axes labels (e.g., xlabel="Time", ylabel="Effort").

With this in mind, create a scatter plot (kind="scatter") of Y and a function of X, give the plot a title ("X vs. Y" is fine) and label each axis ("X", "Y"). You'll need to use the data object and explicitly set the x and y values for the plot (e.g., df.plot(x=, y=, \square)).

² For more information, type help(pd.DataFrame) to get more information on the DataFrame object. Alternately, in the Jupyter notebook, you can press Shift + Tab at the end of a command to get help on the object (i.e., when your cursor is at the end of pd. DataFrame or your new object data.)

³ They only display the results of intermediate lines of a cell if the print() command is used.

⁴ n.b.: Pandas is designed first as a tool for finance and business operations. The default settings for plots are often inappropriate for scientific plots, so some tweaking is often necessary.

Polishing Plot Appearance

When creating more complex plots, once the basics are complete, remove any unnecessary "chart junk" to prepare your plot for publication. This includes things like background shading (Matplotlib is pretty good about this), grid lines (only necessary if you're trying to manually read data off the chart; we don't need this with the ubiquity of computers), and the border surrounding the full plot area (should only need axes to indicate the origin or provide an anchor for tick marks). If there is a legend (useful when there are multiple datasets present), it should usually be a part of the plot area, not pushed off to the side. Don't waste valuable data area in your papers on something like a legend than can be nestled in one of the empty parts of your plot. Finally, while color seems like a nice feature, plots shouldn't rely on color to convey meaning. There's always a chance that readers won't have access to color printouts, or that the reader may be colorblind. To keep plots accessible, use symbols, line weights, and dashing to differentiate your datasets. The lesson to learn from this paragraph is that simplicity is key to conveying your message. The sole purpose of your figures is to help the reader understand the message. Anything that distracts from that is harmful to your cause.

Now that the plot has been cleaned up, we'd like to add a trendline. The Seaborn library is a visual analysis package built on top of Matplotlib with an emphasis on data exploration and friendly figures. To plot our data with Seaborn, input the following:

```
sns.lmplot(data=data, x="X", y="Y", ci=None)
```

This creates a scatterplot using the X and Y columns from data, fits it with a linear model ("lm"), and disables the confidence interval (CI) in the plot.⁵ A linear fit should be sufficient for this data. The Seaborn library is really designed for exploratory data analysis, not for complex curve fits and edge cases of data. It is generally inappropriate for spectral data, but very good for visual and statistical explorations of data.

Fitting Data

Now that you have a trendline, it's time to get some information about it. The NumPy library is a fundamental library in the numeric python community. It contains a vast array of functions and structures that are used by many other libraries. One useful function in NumPy is the Polynomial.fit() function, which calculates a polynomial of best fit using a linear least-squares method. Examples can be seen on the

⁵ "ci" can be set to a numeric value to represent the confidence interval of the associated fit, typically 95% (Ci=95).

⁶ The NumPy library (shortened to np by convention) has a host of useful functions, including a number of common mathematical functions. The square root function is one of these (np.sqrt(num)). Another useful one (that you'll need later) is np.log10(num).

linked page, but we will run through the simplest example, a straight line.

The fit() function is really a *method* of the Polynomial class, meaning it is a function attached to a specific type of object (a Polynomial object). To access it, it's easiest if we import just the Polynomial class with import numpy.polynomial.Polynomial as P. To use the P.fit() function, you need to specify the data ("X" and "Y") and the degree of the polynomial (1). All other arguments are optional, and just giving these three arguments will return a polynomial object ordered from lowest power to highest $(c_0 * x^0, c_1 * x^1, \dots, c_{n-1} * x^{n-1}, c_n * x^n)$. The optional argument full=True will return the full fit statistics as [resid, rank, sv, rcond], including the *covariance matrix*, the diagonal values of which are the variances for each coefficient.

These (or, more precisely, their square roots) help you to see how "good" the returned fit is. Please read the help page to learn what statistics are returned and in what form.⁷

Implement this using the following code:⁸

```
# Fitting using the `numpy.polyfit()` function
   # Function returns fit coefficients (1d array of

    values) and an n x n covariance matrix

   fit coeff, fit cov = np.polyfit(data["X"], data["Y"],

    1, cov=True)

4
   # The standard error of each coefficient is just the
    square root of that diagonal element.
   fit err = np.sqrt(np.diag(fit cov))
   # Use formatted strings (f-strings) to display values

    with useful labels

   print(f"slope: {fit_coeff[0]}, std. err.:
    intercept: {fit coeff[1]}, std. err.: {fit err[1]}")
```

⁷ If you're interested in doing more statistical modeling in Python, the Statsmodel package offers a number of descriptive statistics and flexible models.

⁸ A couple of notes on Python features:

- · everything following the number sign (#) on a line is considered a comment, meaning that it isn't evaluated. This is a great way to document your code as you work.
- The "f-strings" feature was introduced in Python 3.6. They are a far more convenient way to included code in printed strings, and result in much more readable code. Use them if you have the option, I won't bother teaching you older methods of formatting strings.
- Line 3 assigns multiple variables at once by using tuple assignment. Don't worry too much about the mechanics of this, just know that it's a handy tool when the right side of your assignment puts out multiple

Anscombe's Quartet

Repeat the process of importing data with the file anscombe.csv.⁹ Name the new dataframe anscombe and display it by calling it alone in the last line of your cell.¹⁰

Now we want to calculate and print a linear fit and corresponding error for each of the four sets of data in anscombe. The columns in anscombe should have the names "x1", "y1", "x2", "y2", etc. For convenience, I've created a small loop for you to use on your data. This ⁹ There are two options here: you will need to change the delimiter to a comma (delim=",") for this file, or, since the default delimiter for pd.read_csv() is a comma, you can omit the delim=", " argument altogether (it is implicitly defined). 10 Just like we did with the dataframe data in the previous section.

allows us to perform repetitive operations without worrying about being consistent in our typing from operation to operation.¹¹

```
11 If it's correct for the first set of data, it
will be correct for each subsequent one
```

```
# We will make a loop to make this less repetitive
    (and reduce our chances of making an error)
    fits = []
2
    for x, y in
     4 (('x1','y1'),('x2','y2'),('x3','y3'),('x4','y4')):
        fit_coeff, fit_cov = np.polyfit(anscombe[x],

    anscombe[y], 1, cov=True)

        fit err = np.sqrt(np.diag(fit cov))
        fits.append([fit_coeff,fit_err])
6
   # This operation makes my point regarding this data
     ← more clear, as you'll see
   with np.printoptions(precision=2):
        print("Fit param., Std. Err.")
10
        for item in fits:
            print(item)
12
```

Clearly, the data sets are different... You can see that from the raw data in the table. So what's going on? To learn more, plot each of the datasets in a separate plot. Discuss your observations with your lab partner, and figure out the best way to deal with information like this.

The key takeaway from this exercise is that you should always plot your data, even if it's only a quick glance on the side of your workspace! This data was taken from an article written by F.J. Anscombe in 1973 arguing for the necessity of graphs in statistical analysis. It is such a valuable dataset that the seaborn library is installed with a copy of the data (sns.load_dataset("anscombe")).

Column Manipulation and Plotting Spectra

Start a new section titled "Plotting Spectra" and import the file ftir data.csv. Print out the top of the dataframe using the command df.head(), where df is the name of your dataframe. Notice that we have data in the form "Wavenumber (cm ^ -1)", "Transmittance (%)".

There is a very simple syntax in Pandas for making a new column:

```
df["new col name"] = func(df["old col name"])
```

In this case, func() just represents the transformation that we are applying to the original column (multiplication, addition, inversion, etc.). Columns in dataframes will apply functions across every item in the column by default (a process called "broadcasting"), so making a lineby-line transformation (e.g., doubling a column) is as easy as df["B"]

= 2 * df["A"]. Using this syntax, create new calculated columns in our dataframe so we can plot the same spectrum as "Wavelength" vs "Absorbance (A)". 12

To get you started converting wavenumber to wavelength, make the following transformation on the "Wavenumber" column¹³:

```
df["Wavelength (??)"] = 1/(df["Wavenumber
 (cm^{-1})"]/10**(-2)) * 10**(6)
```

You should be able to derive this equation quickly on your own...¹⁴ Perform the same steps to take the column for "Transmittance" and transform it into Absorbance values. Remember that Absorbance is $\log_{10}(1/T)$ while Transmittance is I/I_0 .

Once you have created the new columns, create two plots. The first should plot wavenumber agains transmittance, while the second will plot wavelength against absorbance. Because these are spectra (data collected in a single experiment), it is appropriate to plot them as line plots, without plotting the individual points. Make sure that your plot reflects this.

Make sure your plots are labeled in all the appropriate locations and that your spectra look clean and presentable.

Congratulations, you have the basics of plotting and calculation using Jupyter! You are expected to take this information and build upon it as you write reports through the semester. All of these techniques can be translated to other applications, some of which might be more appropriate for the type of research being performed in a research group. You should be able to transfer your basic skills to any other software with ease, and the tips on presentability are universally applicable.

Alternative Software

While Microsoft Excel may be the most common piece of software for plotting, that doesn't mean it's the best. You have seen that Python (by way of Jupyter) is capable of doing just about anything you can imagine, often in only a few lines. In the future, should you find that Jupyter doesn't meet your needs, here is a list of alternative software for creating scientific plots (I've left out general purpose spreadsheet software).

GraphPad Prism (CUNY site license available) Statistical analysis software for Windows and macOS. Capable of repeated analysis on huge datasets of life/social science data. Falls short in spectral analysis tools.

- 12 What are the relevant units for %T and
- 13 Notice that Python requires the use of "**" for exponentiation. This is because the caret character (^) is often used for other commands. This convention is common in programming languages.
- ¹⁴ Hint: label the units and work through the dimensional analysis. Fill in the question marks with the appropriate units.

https://www.graphpad.com

Origin by OriginLab (no site license available) Professional data analysis and plotting software for scientists and engineers. Heavy focus on physical sciences and engineering. Only available on Windows.

https://www.originlab.com

Igor Pro by WaveMetrics (no site license available) Very similar to Origin, but available for macOS and Windows.

https://www.wavemetrics.com/products/igorpro

Mathematica (site license), Matlab (site license), JupyterLab (free) These are really programming environments, though they are designed for use by scientists. Mathematica and Matlab are commercial products, while Jupyter is an open-source option that uses the JUlia, PYThon, and R languages. They all have entry-output interfaces so that live interaction can be done with data. While the learning curve is steep, the payout for learning a programming language will last a lifetime.

http://www.wolfram.com/mathematica/ https://www.mathworks.com/products/matlab.html https://jupyter.org