

M. Hohle:

Physics 77: Introduction to Computational Techniques in Physics





<u>Week</u>	<u>Date</u>	<u>Topic</u>
1	June 12th	Programming Environment & UIs for Python, Programming Fundamentals
2	June 19th	Basic Types in Python
3	June 26th	Parsing, Data Processing and File I/O, Visualization
4	July 3rd	Functions, Map & Lambda
5	July 10th	Random Numbers & Probability Distributions, Interpreting Measurements
6	July 17th	Numerical Integration and Differentiation
7	July 24th	Root finding, Interpolation
8	July 31st	Systems of Linear Equations, Ordinary Differential Equations (ODEs)
9	Aug 7th	Stability of ODEs, Examples
10	Aug 14th	Final Project Presentations



Outline

Reading/Writing Files

- Pandas and Data Frames
- Faster Alternatives
- More about .txt

Visualization

- Matplotlib
- Seaborn
- plt, ax, fig





Outline

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- Pandas and Data Frames
- Faster Alternatives
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- Matplotlib
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- plt, ax, fig

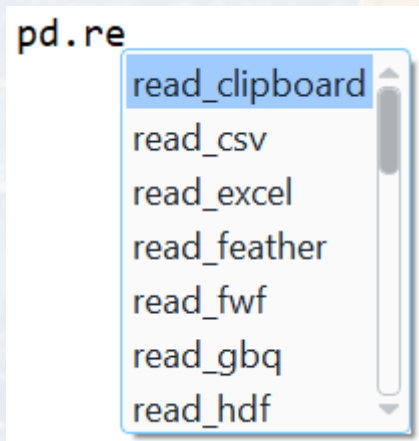




most common file formats

plain text:	.dat .txt .fa
tables:	.csv .xls .xlsx
python:	.py .npy .pkl

```
import pandas as pd
```





most common file formats

plain text: .dat .txt .fa

tables: .csv .xls .xlsx

python: .py .npy .pkl

```
import pandas as pd
```

Microsoft Excel Comma Separated Values File

Data_Set

Molecules

Microsoft Excel Worksheet

Data_Set

Text Document

Cystfibr

```
df = pd.read_excel()
```

11.7 | packaged by
yright", "credits"

.20.0 -- An enhance

```
import pandas as pd
```

```
read_excel(io, sheet_name: 'str | int | list[IntStrT] |  
None'=0, *, header: 'int | Sequence[int] |  
None'=0, names: 'SequenceNotStr[Hashable] | range  
| None'=None, index_col: 'int | str |  
Sequence[int] | None'=None, usecols: 'int | str |  
Sequence[int] | Sequence[str] | Callable[[str],  
bool] | None'=None, dtype: 'DtypeArg |  
None'=None, engine: "Literal['xlrd', 'openpyxl',  
'odf', 'pyxlsb', 'calamine'] | None"=None,  
converters: 'dict[str, Callable] | dict[int,  
Callable] | None'=None, true_values:  
'Iterable[Hashable] | None'=None, false_values:  
'Iterable[Hashable] | None'=None, skiprows:  
'Sequence[int] | int | Callable[[int], object] |  
None'=None, nrows: 'int | None'=None,  
na_values=None, keep_default_na: 'bool'=True,  
na_filter: 'bool'=True, verbose: 'bool'=False,  
parse_dates: 'list | dict | bool'=False,  
date_parser: 'Callable |  
lib.NoDefault'=, date_format:  
'dict[Hashable, str] | str | None'=None,  
thousands: 'str | None'=None, decimal: 'str'='.',  
comment: 'str | None'=None, skipfooter: 'int'=0,  
storage_options: 'StorageOptions | None'=None,  
dtype_backend: 'DtypeBackend |  
lib.NoDefault'=, engine_kwargs: 'dict  
| None'=None)
```

Read an Excel file into a ``pandas`` ``DataFrame``.

Supports `xls`, `xlsx`, `xlsm`, `xlsb`, `odf`, `ods`
and `odt` file extensions read from a local

Parameters

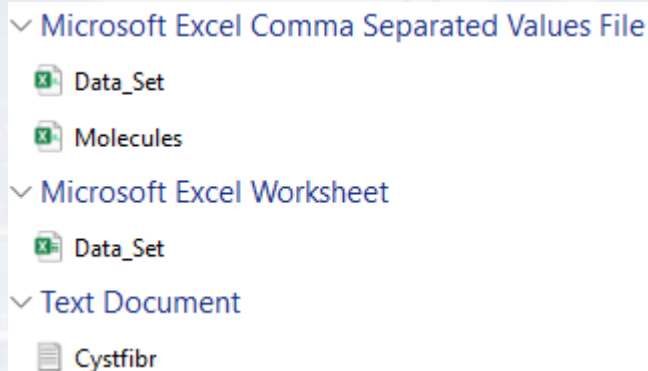
io : str, bytes, ExcelFile, xlrd.Book, path object, or file- ...



most common file formats

plain text:	.dat .txt .fa
tables:	.csv .xls .xlsx
python:	.py .npy .pkl

```
import pandas as pd
```



```
df = pd.read_csv('Molecules.csv')
```

Name	Type	Value
	DataFrame	Column names: molecular_weight, electronegativity, bond_lengths, num_h ...



```
import pandas as pd
```

```
df = pd.read_csv('Molecules.csv')
```

Nam▲	Type	Size	Value
df	DataFrame	(200, 6)	Column names: molecular_weight, electronegativity, bond_lengths, num_h ...

Index	molecular_weight	electronegativity	bond_lengths	num_hydrogen_bonds	logP	label
0	341.704	2.65585	3.09407	2	9.11147	Toxic
1	335.951	3.22262	2.89039	7	8.92848	Toxic
2	235.203	2.44115	2.48203	1	6.49731	Toxic
3	246.505	2.76656	2.71547	7	7.45089	Toxic
4	437.939	3.4801	3.59569	3	10.9156	Toxic
5	336.453	2.81474	3.11	9	8.55696	Toxic
6	372.542	3.17969	3.3866	8	9.48685	Toxic
7	349.19	3.1814	3.19359	7	9.10357	Toxic
8	399.353	3.02359	3.50278	4	9.8421	Toxic



```
import pandas as pd
```

```
df = pd.read_csv('Molecules.csv')
```

```
df.head()
```

Nam▲	Type	Size	Value
df	DataFrame	(200, 6)	Column names: molecular_weight, electronegativity, bond_lengths, num_h ...

```
In [3]: df.head()
```

```
Out[3]:
```

	molecular_weight	electronegativity	...	logP	label
0	341.704142	2.655846	...	9.111473	Toxic
1	335.950798	3.222621	...	8.928483	Toxic
2	235.203185	2.441153	...	6.497307	Toxic
3	246.504930	2.766560	...	7.450888	Toxic
4	437.938926	3.480105	...	10.915629	Toxic

fully equivalent for excel files:

```
df = pd.read_excel('My_File.xlsx')
```



a data frame has numerous functions

`df.head()` shows header

	molecular_weight	electronegativity	...	logP	label
0	341.704142	2.655846	...	9.111473	Toxic
1	335.950798	3.222621	...	8.928483	Toxic
2	235.203185	2.441153	...	6.497307	Toxic

`df.index` returns rows

```
df.index
RangeIndex(start=0, stop=200, step=1)
```

`df.columns` returns columns

```
Index(['molecular_weight', 'electronegativity', 'bond_lengths',  
      'num_hydrogen_bonds', 'logP', 'label'],  
      dtype='object')
```



a data frame has numerous functions

`df.columns` returns columns

```
Index(['molecular_weight', 'electronegativity', 'bond_lengths',  
      'num_hydrogen_bonds', 'logP', 'label'],  
      dtype='object')
```

`df.corr()` returns Pearson's correlation coefficient

columns 0 to 4 contain float/int

```
Corr = df[df.columns[:-1]].corr()
```

Index	molecular_weight	electronegativity	bond_lengths	um_hydrogen_bond	logP
molecular_weight	1	0.0280505	0.953066	0.0157675	0.969772
electronegativity	0.0280505	1	0.0343733	-0.0526109	0.00634745
bond_lengths	0.953066	0.0343733	1	0.0258849	0.926063
num_hydrogen_bonds	0.0157675	-0.0526109	0.0258849	1	0.0104456
logP	0.969772	0.00634745	0.926063	0.0104456	1



a data frame has numerous functions

```
df[['logP', 'label']]
```

returns **data frame** of **selected columns**

	logP	label
0	9.111473	Toxic
1	8.928483	Toxic
2	6.497307	Toxic
3	7.450888	Toxic
4	10.915629	Toxic
..
195	8.794466	Non-Toxic
196	9.651463	Toxic
197	7.651613	Non-Toxic
198	9.060061	Toxic

```
df.loc[[1, 5]]
```

returns **data frame** of **selected rows**

	molecular_weight	electronegativity	...	logP	label
1	335.950798	3.222621	...	8.928483	Toxic
5	336.453422	2.814735	...	8.556958	Toxic



a data frame has numerous functions

Try out the following commands!

```
df.iloc[4:6, 5:9]
```

slicing **data frame** using `iloc`

```
df.iloc[4,5] = 999
```

manipulating individual entries

```
df.insert(2, 'New', df.iloc[:,1])
```

inserting another column called *New*

```
df.rename(index = {1: 'bbb'})
```

changing name of row **1**

```
df.rename(columns = {'Label': 'Toxic or Not'})
```

changing column name



```
import pandas as pd
```

```
df = pd.read_csv('Molecules.csv')
```

finally, saving the data frame to an excel file

```
df.t
```

- tail
- take
- to_clipboard
- to_csv
- to_dict
- to_excel**
- to_feather

```
df.to_excel('Molecules.xlsx')
```

Name	Date Modified
Cystfibr.txt	11/09/2024 04:24
Data_Set.csv	07/12/2023 20:16
Data_Set.xlsx	05/12/2023 20:04
Molecules.csv	05/09/2024 20:30
Molecules.xlsx	06/03/2025 21:30

Microsoft Excel Comma Separated Values File
Data_Set
Molecules
Microsoft Excel Worksheet
Data_Set
Molecules



pandas can also read **text files**:

```
cf = pd.read_csv('cystfibr.txt')
```



cf - DataFrame

Index	age	sex	height	weight	bmp	fev1	rv	frc	tlc	pemax
0	7	0	109	13.1	68	32				
	258	183	137	95						
1	7	1	112	12.9	65	19				
	449	245	134	85						
2	8	0	124	14.1	64	22				
	441	268	147	100						

```
cf = pd.read_csv('cystfibr.txt', sep = '\\s+')
```

cf - DataFrame

Index	age	sex	height	weight	bmp	fev1	rv	frc	tlc	pemax
0	7	0	109	13.1	68	32	258	183	137	95
1	7	1	112	12.9	65	19	449	245	134	85
2	8	0	124	14.1	64	22	441	268	147	100
3	8	1	125	16.2	67	41	234	146	124	85



Outline

Reading/Writing Files

- Pandas and Data Frames
- **Faster Alternatives**
- More about .txt

Visualization

- Matplotlib
- Seaborn
- plt, ax, fig





pandas is the standard library, but it is slow

reading a 130MB **excel file**:

```
import pandas as pd
import time
```

```
t1 = time.monotonic()
```

```
df = pd.read_excel('Data_Set.xlsx')
```

```
t2 = time.monotonic()
```

```
dt = t2 - t1
```

```
print("Total runtime: " + str(dt) + ' seconds')
```

83 seconds



pandas is the standard library, but it is slow

reading a 180MB **csv file** with the **same content**:

```
import pandas as pd  
import time
```

```
t1 = time.monotonic()
```

```
df = pd.read_csv('Data_Set.csv')
```

```
t2 = time.monotonic()
```

```
dt = t2 - t1
```

```
print("Total runtime: " + str(dt) + ' seconds')
```

1.2 seconds!



pandas:

excel file:	83.0 sec
csv file:	1.2 sec

reading a 180MB **txt file** with the **same content**:

```
import pandas as pd
import time
```

```
t1 = time.monotonic()
```

```
df = pd.read_csv('Data_Set.txt')
```

```
t2 = time.monotonic()
```

```
dt = t2 - t1
```

```
print("Total runtime: " + str(dt) + ' seconds')
```

1.5 seconds!



pandas:

excel file:	83.0 sec
csv file:	1.2 sec
txt file:	1.5 sec

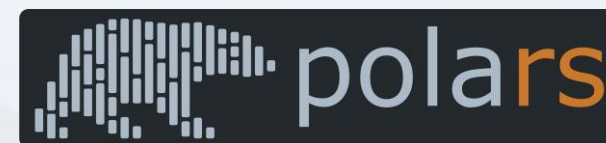


dask



faster than pandas,
but fewer functions

polars



fireducks

FireDucks



pandas:

excel file:	83.0 sec
csv file:	1.2 sec
txt file:	1.5 sec

```
pip install dask  
pip install polars  
pip install xlsw2csv #for excel API  
pip install fastexcel
```



```
import dask.dataframe as dd
```



pandas:

excel file:	83.0 sec
csv file:	1.2 sec
txt file:	1.5 sec

```
import dask.dataframe as dd
```

```
t1 = time.monotonic()
```

```
df = dd.read_csv('Data_Set.csv')
```

```
t2 = time.monotonic()
```

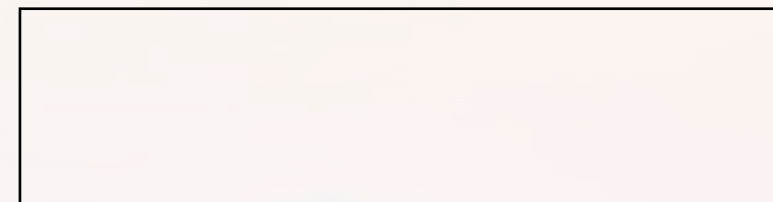
```
dt = t2 - t1
```

```
print("Total runtime: " + str(dt) + ' seconds')
```

0.016 seconds!

```
df = pd.DataFrame(df)
```

However, we might need to transform the output



```
import dask.dataframe as dd
```

```
t1 = time.monotonic()
```

```
df = pd.DataFrame(dd.read_csv('Data_Set.csv'))
```

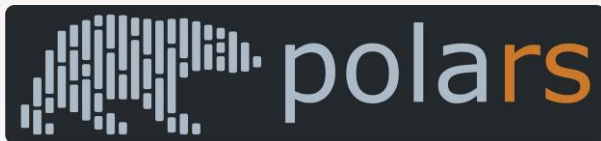
```
t2 = time.monotonic()
```

```
dt = t2 - t1
```

1.7 seconds!

```
print("Total runtime: " + str(dt) + ' seconds')
```

As of Mar 2025, dask **doesn't have an excel API**



```
import polars as pl
```

```
t1 = time.monotonic()
```

```
df = pl.read_excel('Data_Set.xlsx')
```

```
t2 = time.monotonic()
```

```
dt = t2 - t1
```

```
print("Total runtime: " + str(dt) + ' seconds')
```

```
df = pd.DataFrame(df)
```

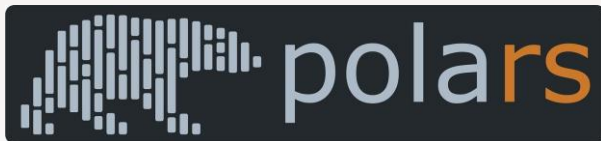
pandas:

excel file:	83.0	sec
csv file:	1.2	sec
txt file:	1.5	sec

dask	csv file:	0.016	sec
	to df	1.7	sec

8.2 seconds!

However, we might need to transform the output



```
import polars as pl
```

```
t1 = time.monotonic()
```

```
df = pd.DataFrame(pl.read_excel('Data_Set.xlsx'))
```

```
t2 = time.monotonic()
```

```
dt = t2 - t1
```

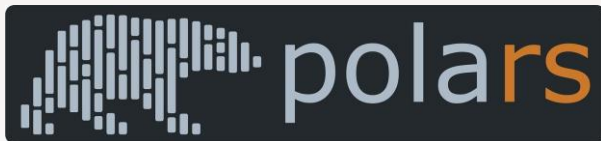
```
print("Total runtime: " + str(dt) + ' seconds')
```

pandas:

excel file:	83.0	sec
csv file:	1.2	sec
txt file:	1.5	sec

dask	csv file:	0.016	sec
	to df	1.7	sec

10.2 seconds!



```
import polars as pl
```

```
t1 = time.monotonic()
```

```
df = pl.read_csv('Data_Set.csv')
```

```
t2 = time.monotonic()
```

```
dt = t2 - t1
```

```
print("Total runtime: " + str(dt) + ' seconds')
```

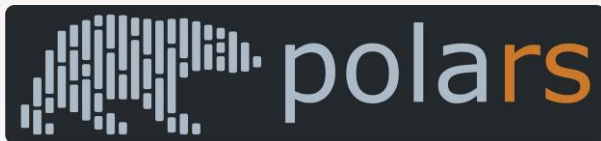
pandas:

excel file:	83.0	sec
csv file:	1.2	sec
txt file:	1.5	sec

dask	csv file:	0.016	sec
	to df	1.7	sec

polars	excel file	8.2	sec
	to df	10.2	sec

0.27 seconds!



```
import polars as pl
```

```
t1 = time.monotonic()
```

```
df = pd.DataFrame(pl.read_csv('Data_Set.csv'))
```

```
t2 = time.monotonic()
```

```
dt = t2 - t1
```

```
print("Total runtime: " + str(dt) + ' seconds')
```

pandas:	excel file:	83.0	sec
	csv file:	1.2	sec
	txt file:	1.5	sec
dask	csv file:	0.016	sec
	to df	1.7	sec
polars	excel file	8.2	sec
	to df	10.2	sec

0.20 seconds!



.xlsx

83 sec

na

8.2 sec
10.2 sec

.csv

1.2 sec

0.016 sec
1.6 sec

0.27 sec
0.20 sec

.txt

1.5 sec

0.016 sec
0.92 sec

0.25 sec
0.23 sec



.xlsx

83 sec

na

8.2 sec

10.2 sec

.csv

1.2 sec

0.016 sec

1.6 sec

0.27 sec

0.20 sec

.txt

1.5 sec

0.016 sec

0.92 sec

0.25 sec

0.23 sec



Outline

Reading/Writing Files

- Pandas and Data Frames
- Faster Alternatives
- **More about .txt**

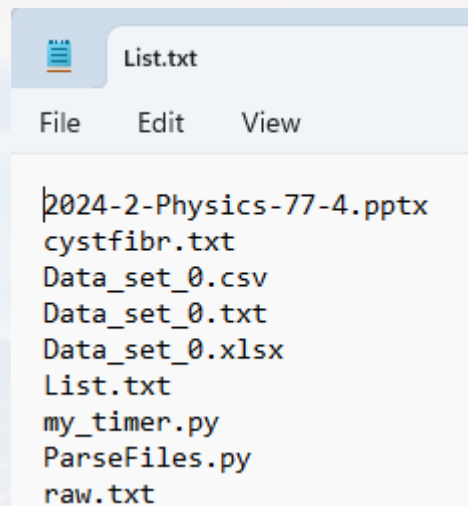
Visualization

- Matplotlib
- Seaborn
- plt, ax, fig





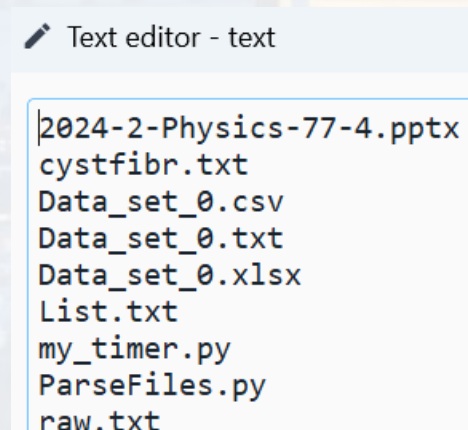
sometimes txt files don't come in a nice format



→ using **open**

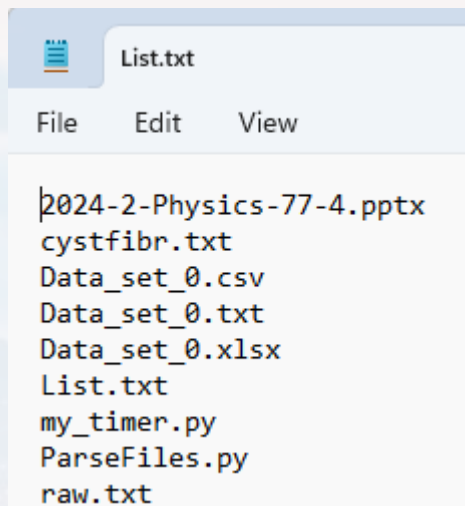
```
with open('List.txt', errors = "ignore") as f:  
    text = f.read()
```

reads file line by line and stops automatically when has reached the end

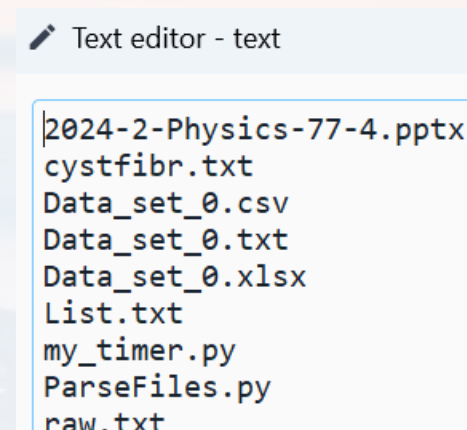




sometimes txt files don't come in a nice format



```
with open('List.txt', errors = "ignore") as f:  
    text = f.read()
```



```
T = list(text.split())
```

Index ▲	Type	Size	
0	str	24	2024-2-Physics-77-4.pptx
1	str	12	cystfibr.txt
2	str	14	Data_set_0.csv
3	str	14	Data_set_0.txt
4	str	15	Data_set_0.xlsx
5	str	8	List.txt
6	str	11	my_timer.py



syntax of **open**

```
with open('my_file.txt', flags, errors = "ignore") as f:  
    text = f.read()
```

finishes once it has reached the end
and closes *f*

flags/modes:

'r'	opens file for reading only.
'w'	opens file for writing. If the file exists, it overwrites it, otherwise, it creates a new file.
'a'	opens file for appending only. If the file doesn't exist, it creates the file.
'x'	creates new file. If the file exists, it fails.
'+'	opens file for updating
't'	opens file in text mode (default)
'b'	opens file in binary mode

flags can be combined like, eg 'wb'

with opens a loop for reading the file line by line



syntax of **open**

```
with open('List.txt', 'r') as read_f:
```

opens file line by line

```
with open('List_copy.txt', 'w') as write_f:
```

```
for r in read_f:  
    write_f.write(r)
```

opens **new** file line by line
for **writing**

writes each line to **new** file

```
In [5]: print(r)  
raw.txt
```

```
In [6]: print(type(r))  
<class 'str'>
```



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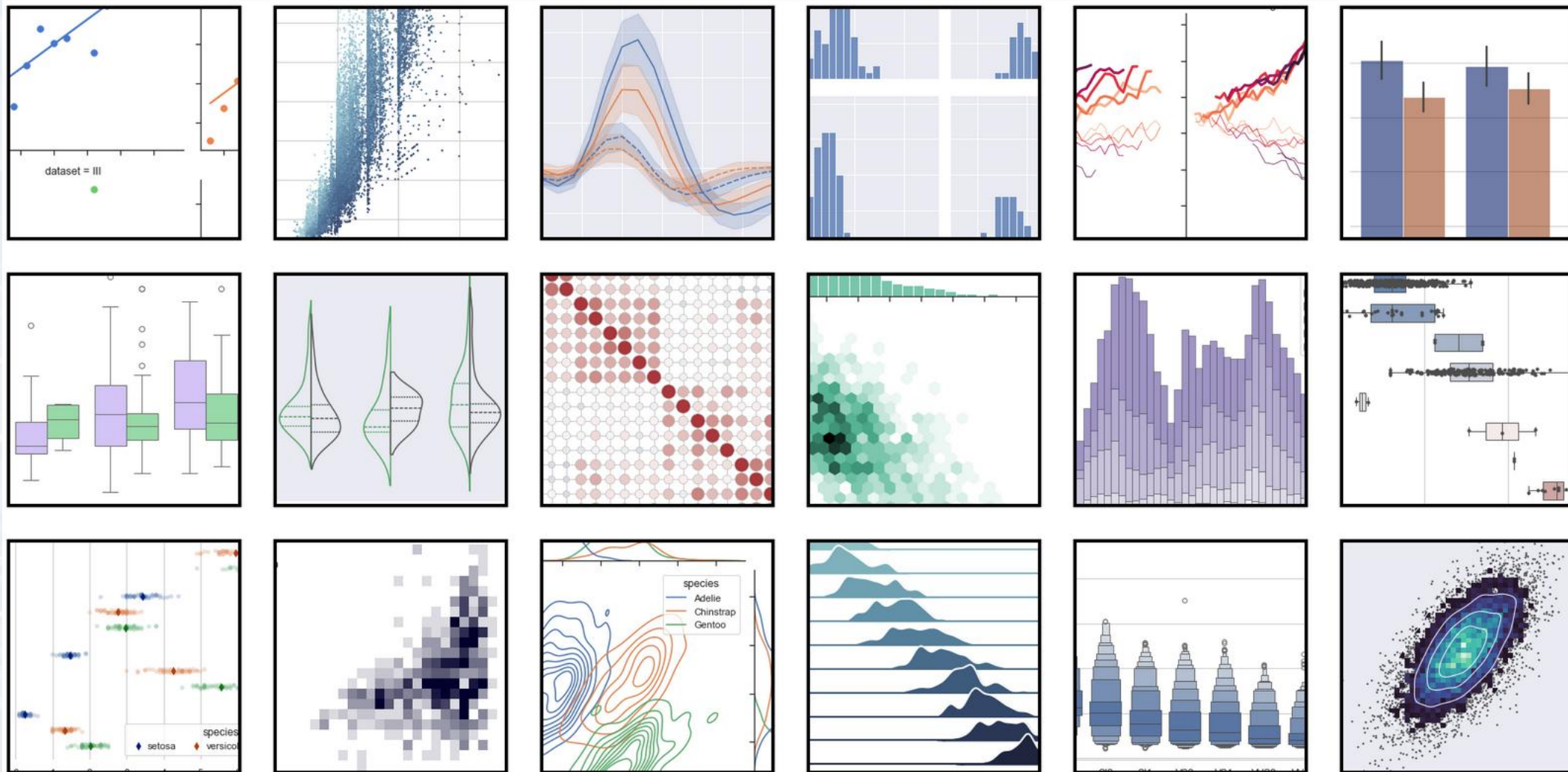
Visualization

- **Matplotlib**
- Seaborn
- plt, ax, fig





Python Graph Gallery





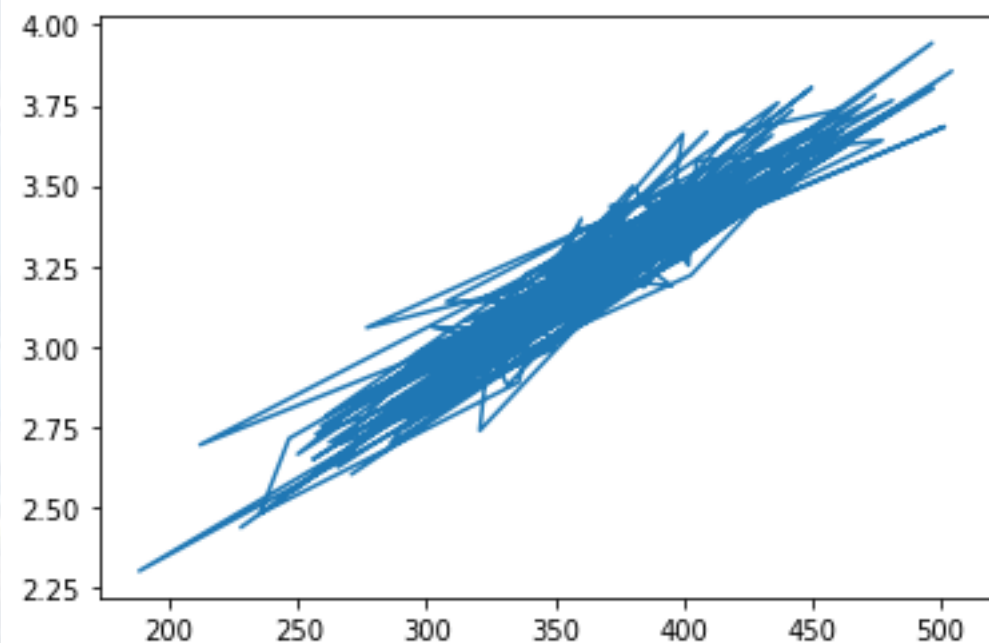
```
import pandas as pd  
import matplotlib.pyplot as plt
```

```
Data = pd.read_csv('Molecules.csv')
```

```
x = Data.molecular_weight  
y = Data.bond_lengths
```

```
plt.plot(x,y)
```

basic plots
arguments
settings





```
import pandas as pd  
import matplotlib.pyplot as plt
```

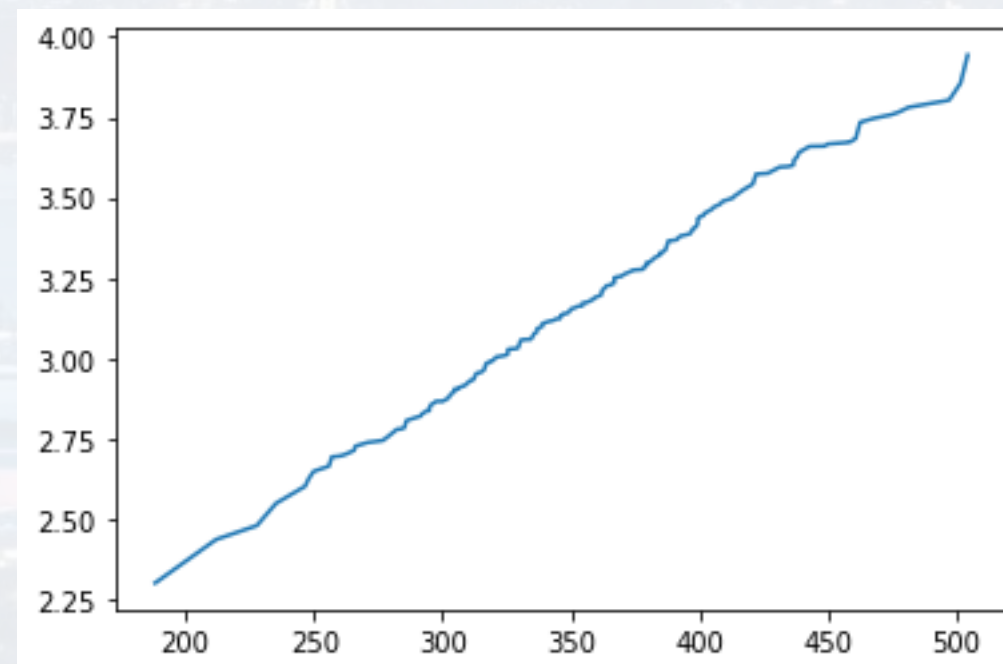
```
Data = pd.read_csv('Molecules.csv')
```

```
x = Data.molecular_weight  
y = Data.bond_lengths
```

```
plt.plot(sorted(x), sorted(y))
```

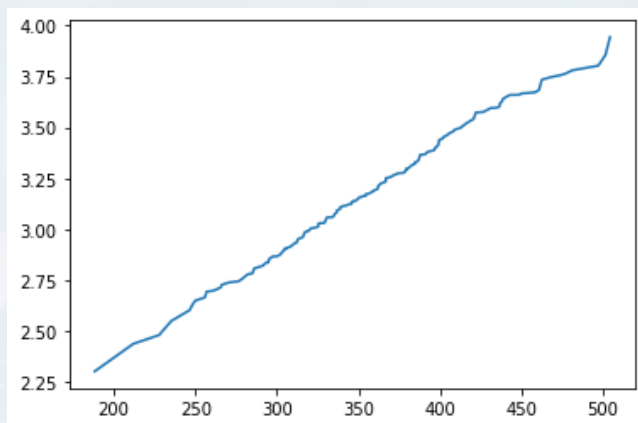
basic plots

arguments
settings

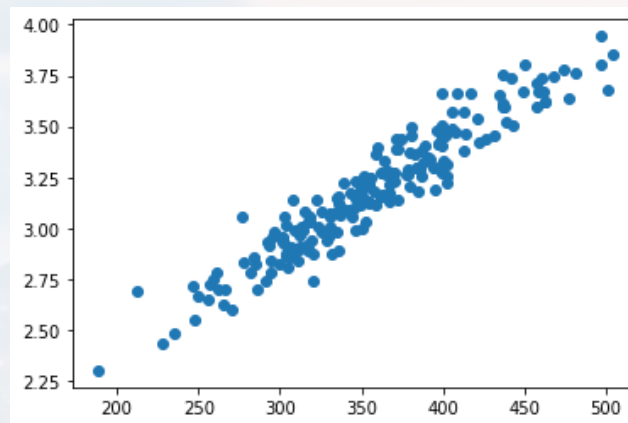




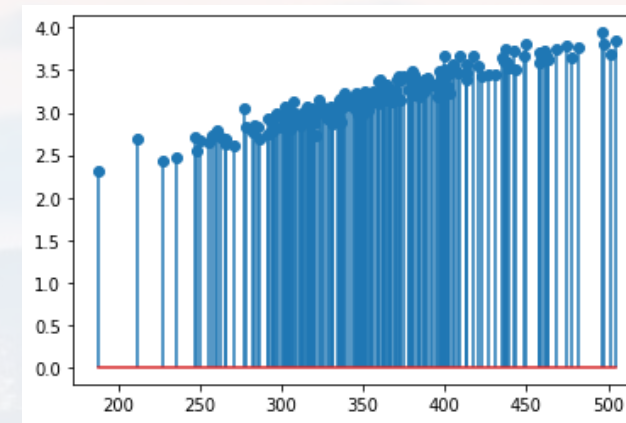
```
plt.plot(sorted(x),  
         sorted(y))
```



```
plt.scatter(x, y)
```



```
plt.stem(x, y)
```

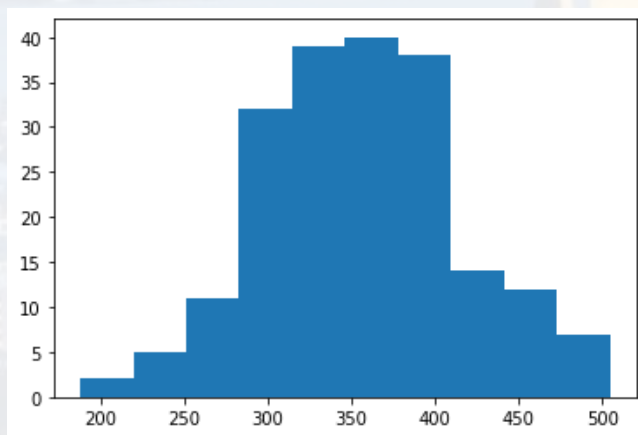


basic plots

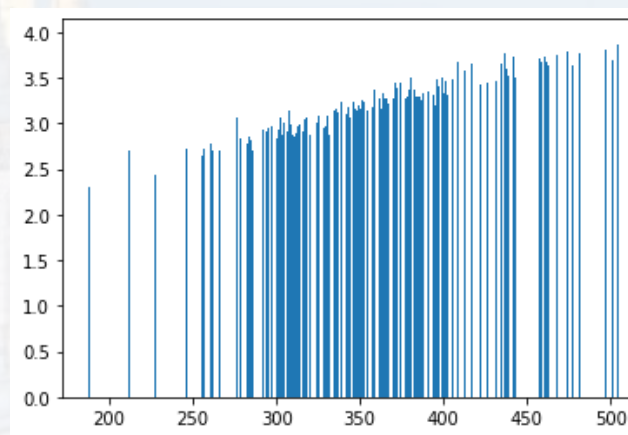
arguments

settings

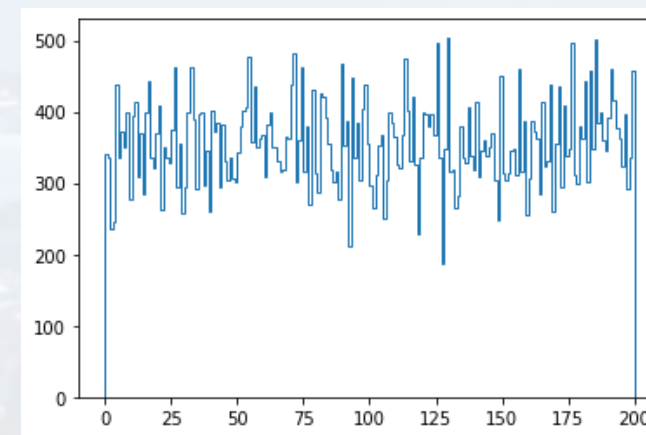
```
plt.hist(x)
```



```
plt.bar(x, y)
```



```
plt.stairs(x)
```





`plt.scatter(`

```
scatter(x: 'float | ArrayLike', y: 'float | ArrayLike', s:  
        'float | ArrayLike | None' = None, c: 'ArrayLike |  
        Sequence[ColorType] | ColorType | None' = None,  
        marker: 'MarkerType | None' = None, cmap: 'str |  
        Colormap | None' = None, norm: 'str | Normalize |  
        None' = None, vmin: 'float | None' = None, vmax:  
        'float | None' = None, alpha: 'float | None' = None,  
        linewidths: 'float | Sequence[float] | None' = None,  
        *, edgecolors: "Literal['face', 'none'] | ColorType  
        | Sequence[ColorType] | None" = None, plotnonfinite:  
        'bool' = False, data=None, **kwargs,)
```

A scatter plot of *y* vs. *x* with varying marker size and/or color.

Parameters

x, *y* : float or array-like, shape (n,)

The data positions. ...



```
plt.scatter(x, y,  
            marker = 'p',  
            s = 135,  
            color = [0.5, 0.1, 0.1],  
            edgecolor = 'black',  
            alpha = 0.3)
```

marker style: **str**

basic plots
arguments
settings

marker	symbol	description
"."	•	point
"."	.	pixel
"o"	●	circle
"v"	▼	triangle_down
"^"	▲	triangle_up
"<"	◀	triangle_left
">"	▶	triangle_right
"1"	⋿	tri_down
"2"	⋿	tri_up
"3"	⋿	tri_left
"4"	⋿	tri_right
"8"	◼	octagon
"s"	◼	square
"p"	⬠	pentagon
"D"	⬠	plus (filled)
"x"	✱	star
"h"	⬠	hexagon1
"H"	⬠	hexagon2

"+"	+	plus
"x"	×	x
"X"	✱	x (filled)
"D"	◆	diamond
"d"	◆	thin_diamond
" "		vline
"_"	—	hline
0 (TICKLEFT)	—	tickleft
1 (TICKRIGHT)	—	tickright
2 (TICKUP)		tickup
3 (TICKDOWN)		tickdown
4 (CARETLEFT)	◀	caretleft
5 (CARETRIGHT)	▶	caretright
6 (CARETUP)	▲	caretup
7 (CARETDOWN)	▼	caredown
8 (CARETLEFTBASE)	◀	caretleft (centered at base)
9 (CARETRIGHTBASE)	▶	caretright (centered at base)
10 (CARETUPBASE)	▲	caretup (centered at base)
11	▼	caredown (centered at base)



basic plots
arguments
settings

```
plt.scatter(x, y,  
            marker = 'p',  
            s = 135,  
            color = [0.5, 0.1, 0.1],  
            edgecolor = 'black',  
            alpha = 0.3)
```

marker size in
pixel: **int**



basic plots
arguments
settings

```
plt.scatter(x, y,  
            marker = 'p',  
            s = 135,  
            color = [0.5, 0.1, 0.1],  
            edgecolor = 'black',  
            alpha = 0.3)
```

color:

array RGB code if **three** values, RGB code plus alpha, if **four** values

str	full string:	'green' 'yellow'
	abbreviation:	'g' 'y'
	HEX code:	'#4b8333' '#fff8de'



basic plots
arguments
settings

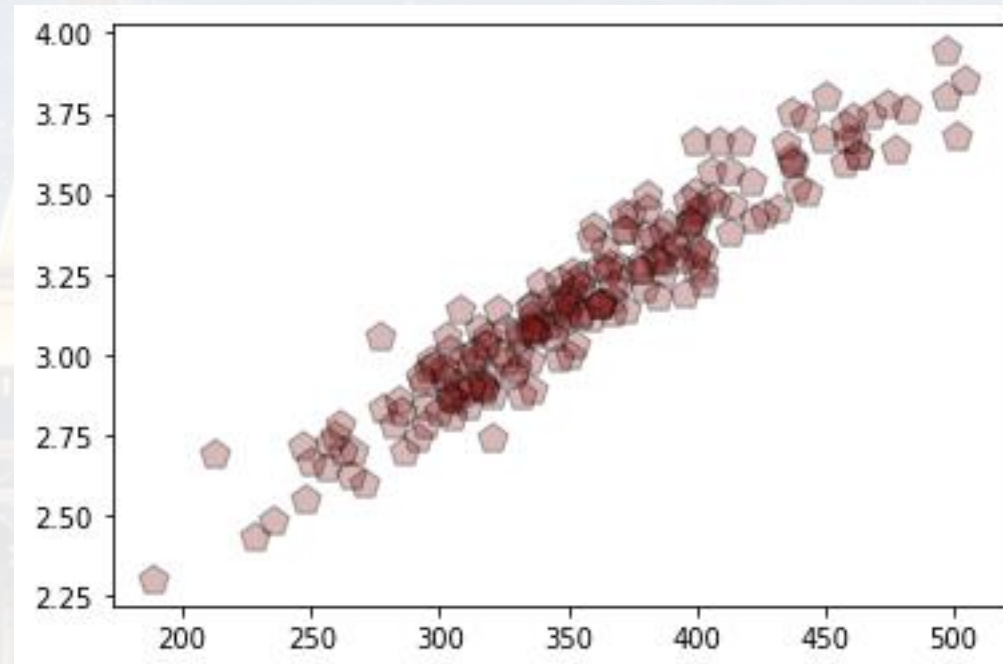
```
plt.scatter(x, y,  
            marker = 'p',  
            s = 135,  
            color = [0.5, 0.1, 0.1],  
            edgecolor = 'black',  
            alpha = 0.3)
```

alpha (opaqueness/opacity): float



basic plots
arguments
settings

```
plt.scatter(x, y,  
            marker = 'p',  
            s = 135,  
            color = [0.5, 0.1, 0.1],  
            edgecolor = 'black',  
            alpha = 0.3)
```





```
plt.scatter(x, y, marker = 'p', s = 135, color = [0.5, 0.1, 0.1],  
           edgecolor = 'black', alpha = 0.3)
```

basic plots
arguments
settings

```
plt.xlabel(r'x values  $\tau^{ij}_{def}$ ')  
plt.ylabel('y values')  
plt.title('first plot')  
plt.legend(['data'])  
plt.xscale('log')  
plt.savefig('new_plot.pdf')  
plt.show()
```

Python speaks LaTeX, but needs
raw string



```
plt.scatter(x, y, marker = 'p', s = 135, color = [0.5, 0.1, 0.1],  
           edgecolor = 'black', alpha = 0.3)
```

basic plots
arguments
settings

```
plt.xlabel(r'x values  $\tau^{ij}_{def}$ ')  
plt.ylabel('y values')  
plt.title('first plot')  
plt.legend(['data'])  
plt.xscale('log')  
plt.savefig('new_plot.pdf')  
plt.show()
```

legend needs to be
a list



```
plt.scatter(x, y, marker = 'p', s = 135, color = [0.5, 0.1, 0.1],  
           edgecolor = 'black', alpha = 0.3)
```

basic plots
arguments
settings

```
plt.xlabel(r'x values  $\tau^{ij}_{def}$ ')  
plt.ylabel('y values')  
plt.title('first plot')  
plt.legend(['data'])  
plt.xscale('log')  
plt.savefig('new_plot.pdf')  
plt.show()
```

plots can be saved
to any common
format



```
plt.scatter(x, y, marker = 'p', s = 135, color = [0.5, 0.1, 0.1],  
           edgecolor = 'black', alpha = 0.3)
```

basic plots
arguments
settings

```
plt.xlabel(r'x values  $\tau^{ij}_{def}$ ')  
plt.ylabel('y values')  
plt.title('first plot')  
plt.legend(['data'])  
plt.xscale('log')  
plt.savefig('new_plot.pdf')  
plt.show()
```

sometimes plots don't show up
(depending on settings)
type `plt.show()`
at the **very end**

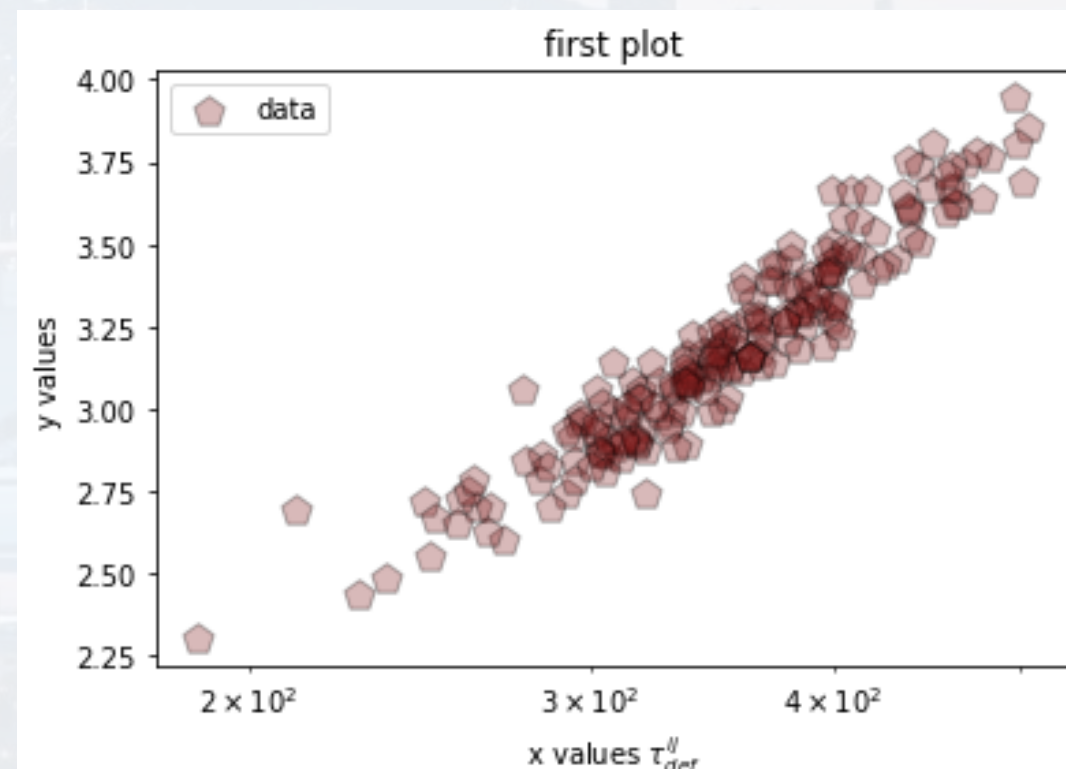


```
plt.scatter(x, y, marker = 'p', s = 135, color = [0.5, 0.1, 0.1],  
           edgecolor = 'black', alpha = 0.3)
```

basic plots
arguments
settings

```
plt.xlabel(r'x values  $\tau_{der}^{ij}$ ')  
plt.ylabel('y values')  
plt.title('first plot')  
plt.legend(['data'])  
plt.xscale('log')  
plt.savefig('new_plot.pdf')  
plt.show()
```

Name	Date Modified
Cystfibr.txt	11/09/2024 04:24
Data_Set.csv	07/12/2023 20:16
Data_Set.txt	16/09/2024 22:37
Data_Set.xlsx	05/12/2023 20:04
List.txt	16/09/2024 23:26
Molecules.csv	05/09/2024 20:30
Molecules.xlsx	06/03/2025 21:30
new_plot.pdf	07/03/2025 01:17





more than one data set:

basic plots
arguments
settings

```
for a in range(3):  
    a += 1  
    plt.scatter(x, y**a,  
                marker = 'p',  
                s       = 30,  
                alpha   = 0.3,  
                label   = 'Data Set ' + str(a))
```

```
plt.xlabel(r'x values')  
plt.ylabel('y values')  
plt.title('first plot')  
plt.legend()  
plt.show()
```

assigns a label to each data
set → stored for legend

calling the legend (no input
argument)

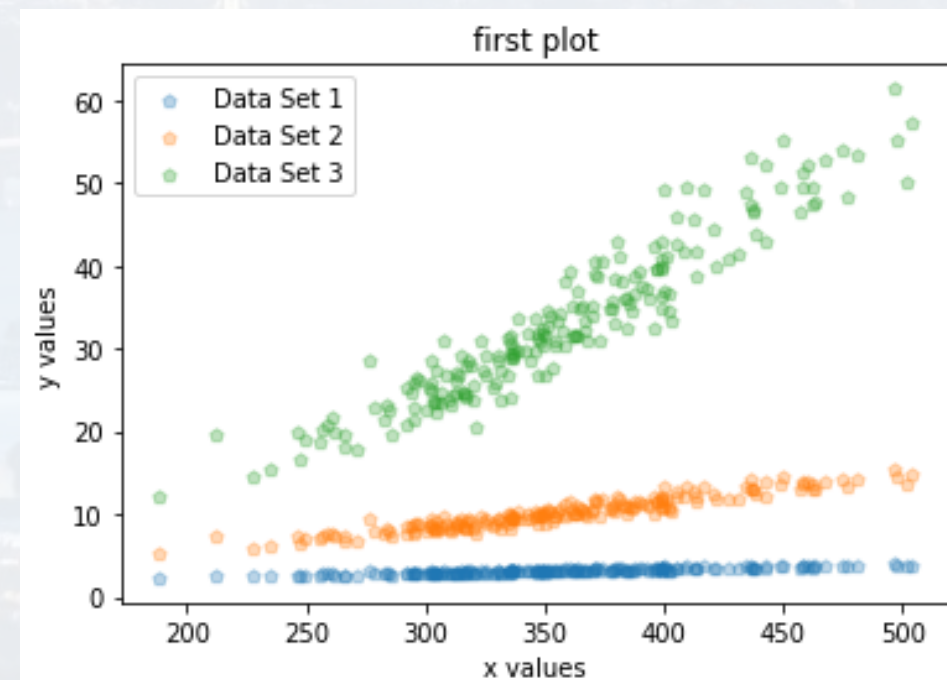


more than one data set:

```
for a in range(3):  
    a += 1  
    plt.scatter(x, y**a,  
                marker = 'p',  
                s       = 30,  
                alpha   = 0.3,  
                label   = 'Data Set ' + str(a))
```

```
plt.xlabel(r'x values')  
plt.ylabel('y values')  
plt.title('first plot')  
plt.legend()  
plt.show()
```

basic plots
arguments
settings





Outline

Reading/Writing Files

- Pandas and Data Frames
- Faster Alternatives
- More about .txt

Visualization

- Matplotlib
- **Seaborn**
- plt, ax, fig



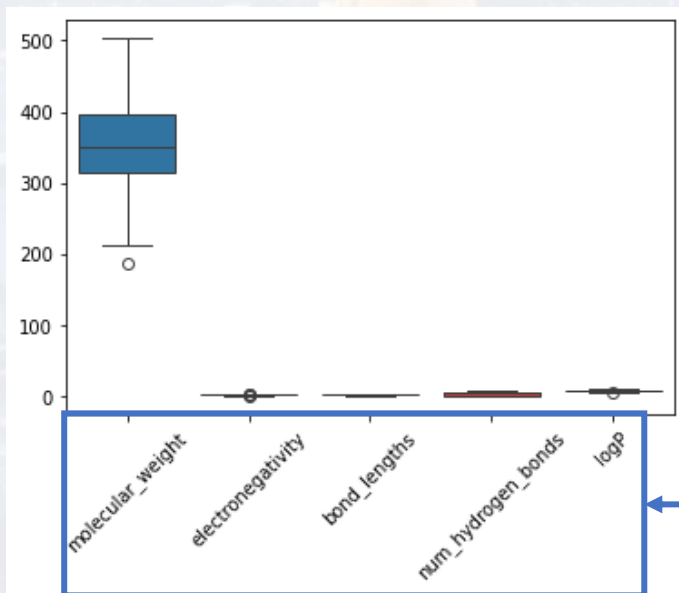


sophisticated plots:

```
import seaborn as sns
```

starting from **data frames** right away

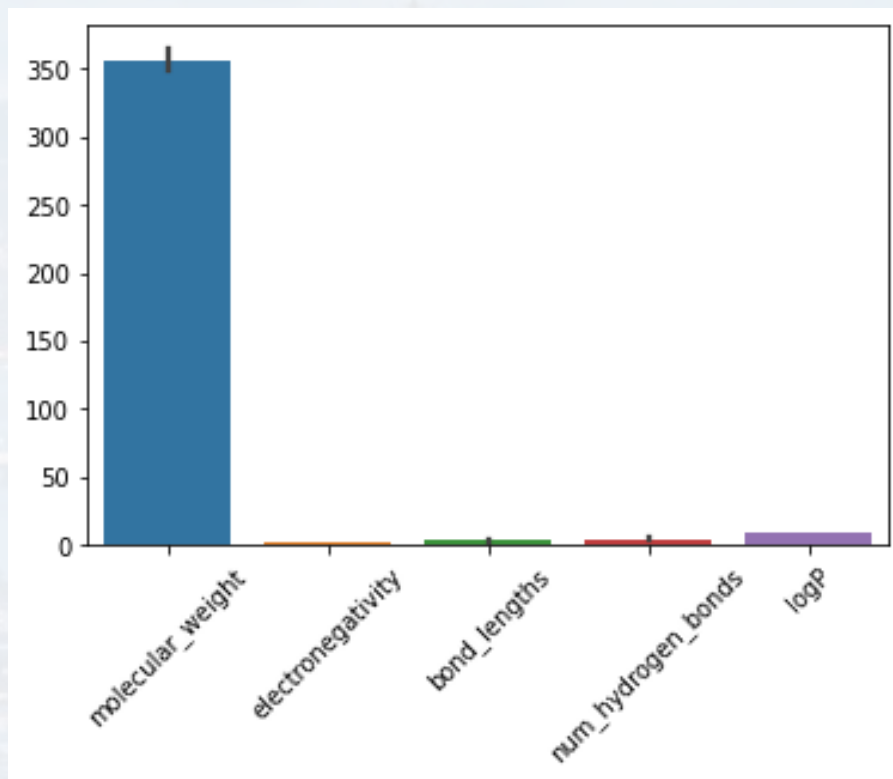
```
sns.boxplot(Data)  
plt.xticks(rotation = 45)  
plt.show()
```



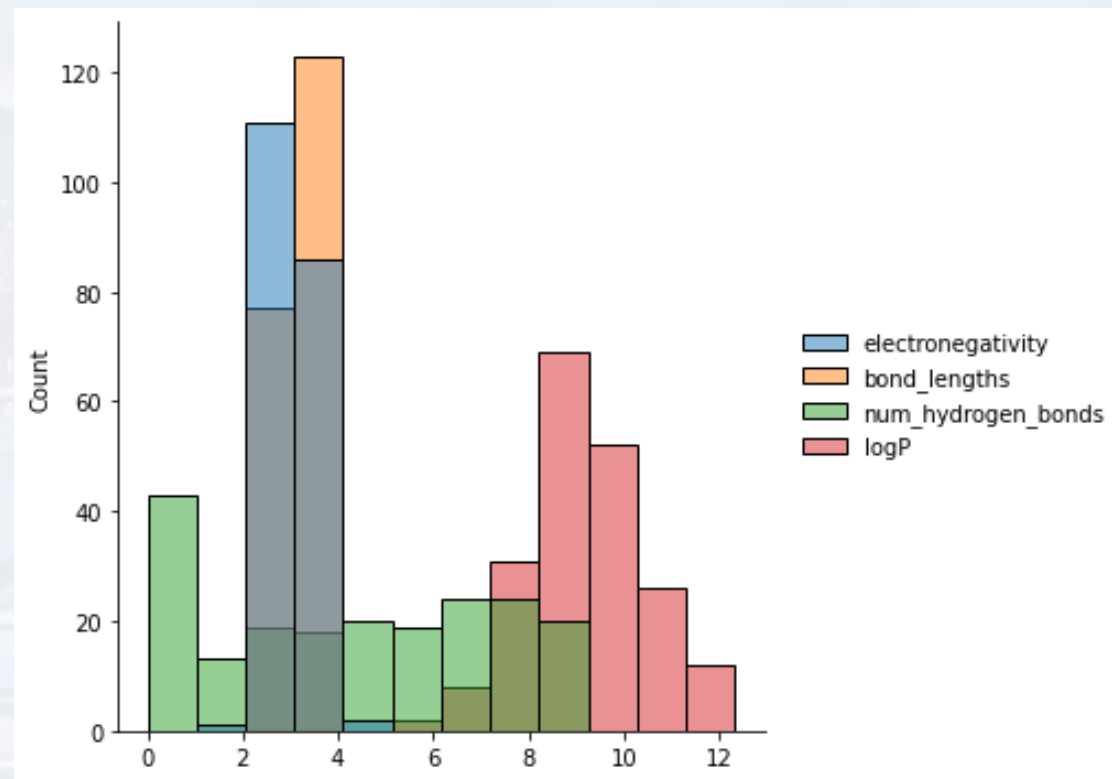
reads labels from data
frame



```
sns.barplot(Data)  
plt.xticks(rotation = 45)  
plt.show()
```

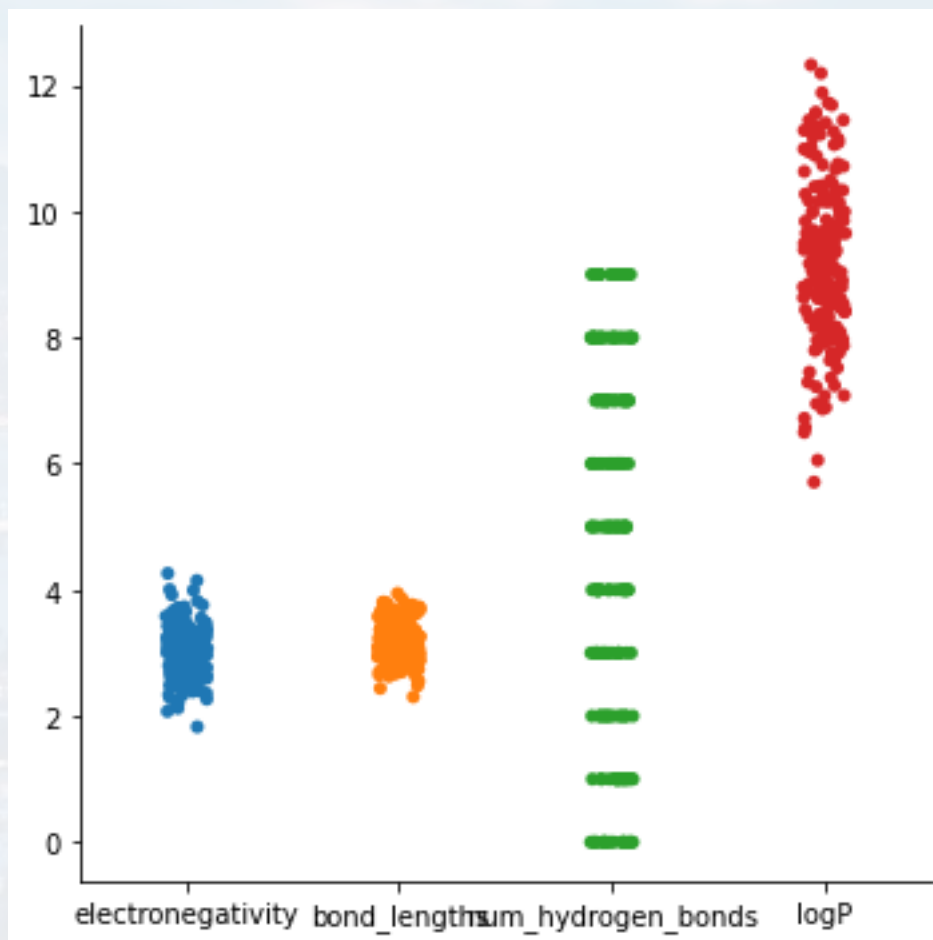


```
sns.displot(Data[Data.columns[1:-1]])  
plt.show()
```



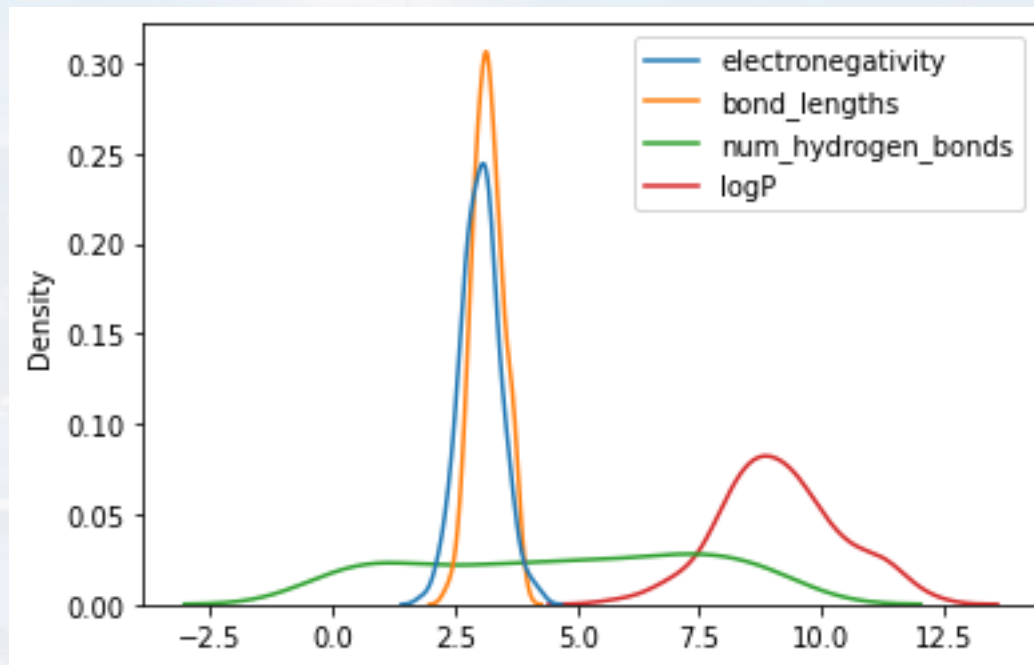


```
sns.catplot(Data[Data.columns[1:-1]])  
plt.show()
```





```
sns.kdeplot(Data[Data.columns[1:-1]])  
plt.show()
```



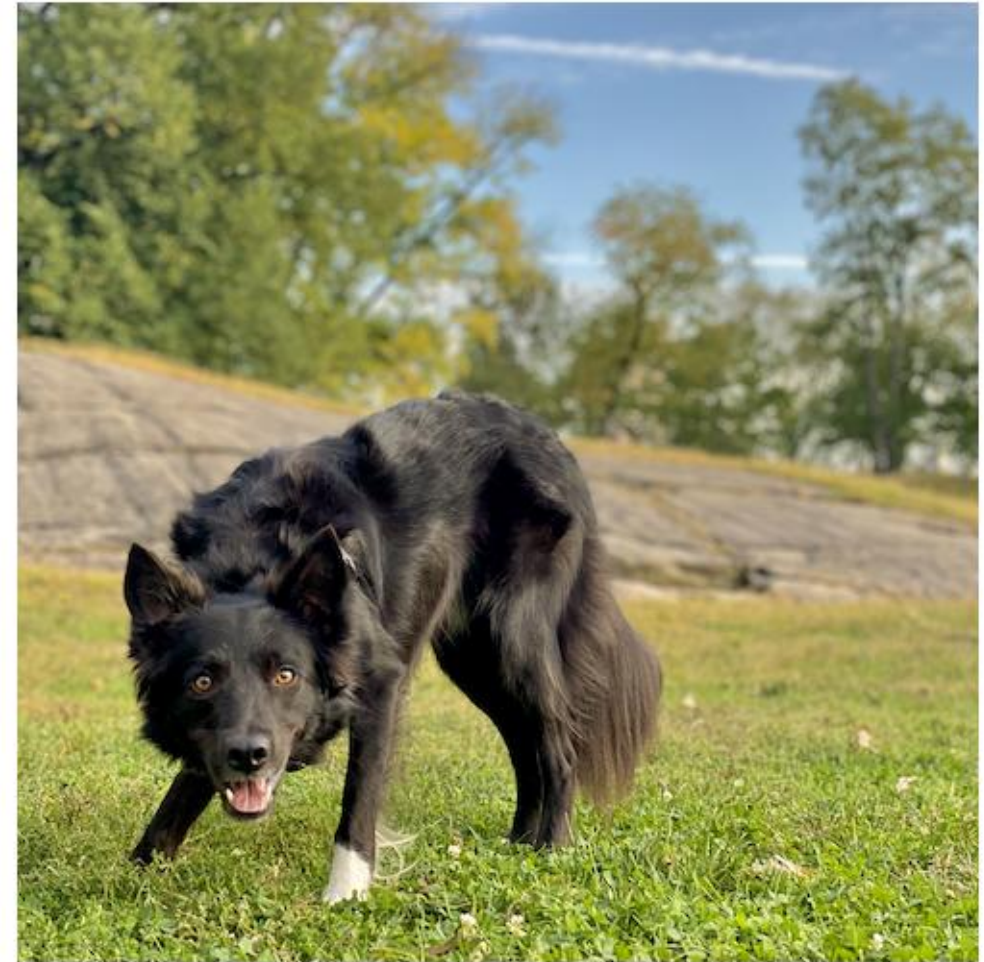
[Graph Gallery with examples](#)



`sns.dogplot()`



`sns.dogplot(1)`





Outline

Reading/Writing Files

- Pandas and Data Frames
- Faster Alternatives
- More about .txt

Visualization

- Matplotlib
- Seaborn
- **plt, ax, fig**





Three levels in Python:

plt

the plot itself

ax

referring to axis of specific plot

fig

referring to a figure (which can include numerous subplots)

mosaic subplots

classical subplots





Three levels in Python:

plt

the plot itself

ax

referring to axis of specific plot

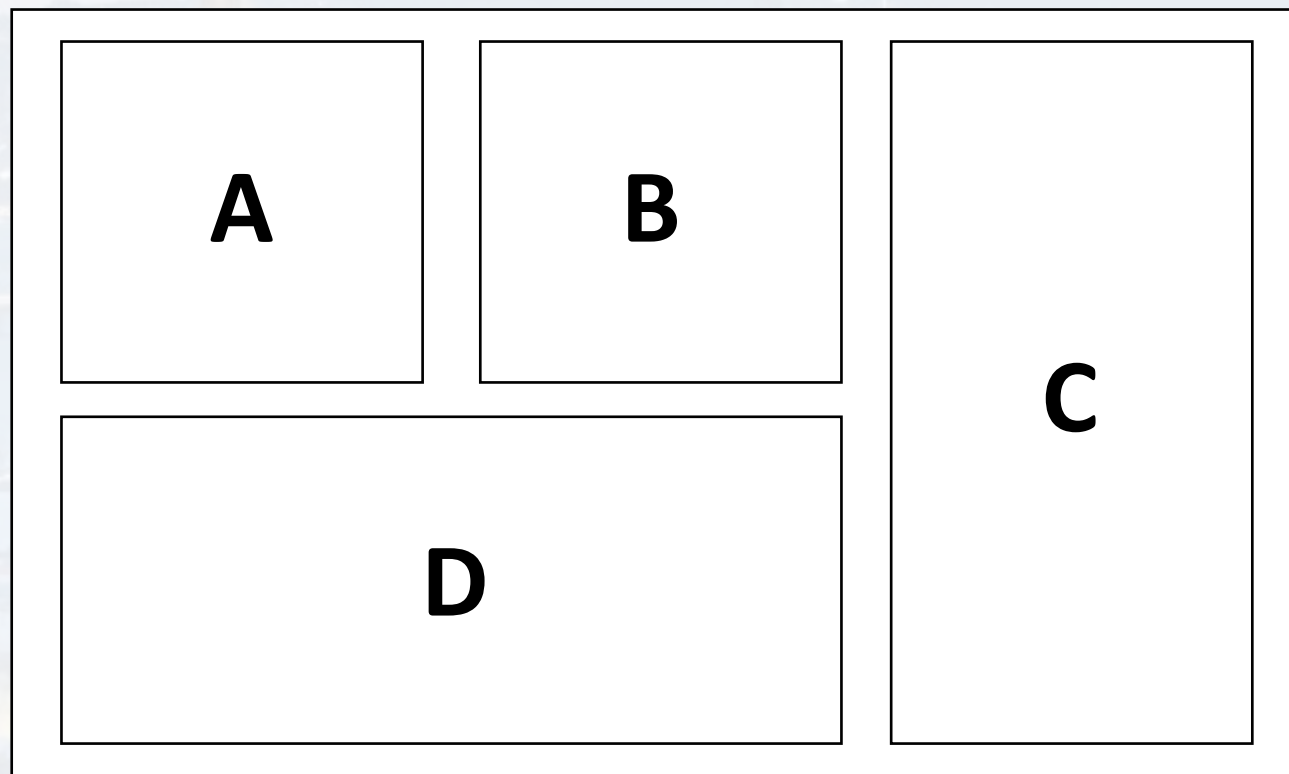
fig

referring to a figure (which can include numerous subplots)

mosaic subplots

classical subplots

```
figMo, axes = plt.subplot_mosaic([[ 'A', 'B', 'C' ],\n                                   [ 'D', 'D', 'C' ]], layout = "constrained")
```





Three levels in Python:

plt

the plot itself

ax

referring to axis of specific plot

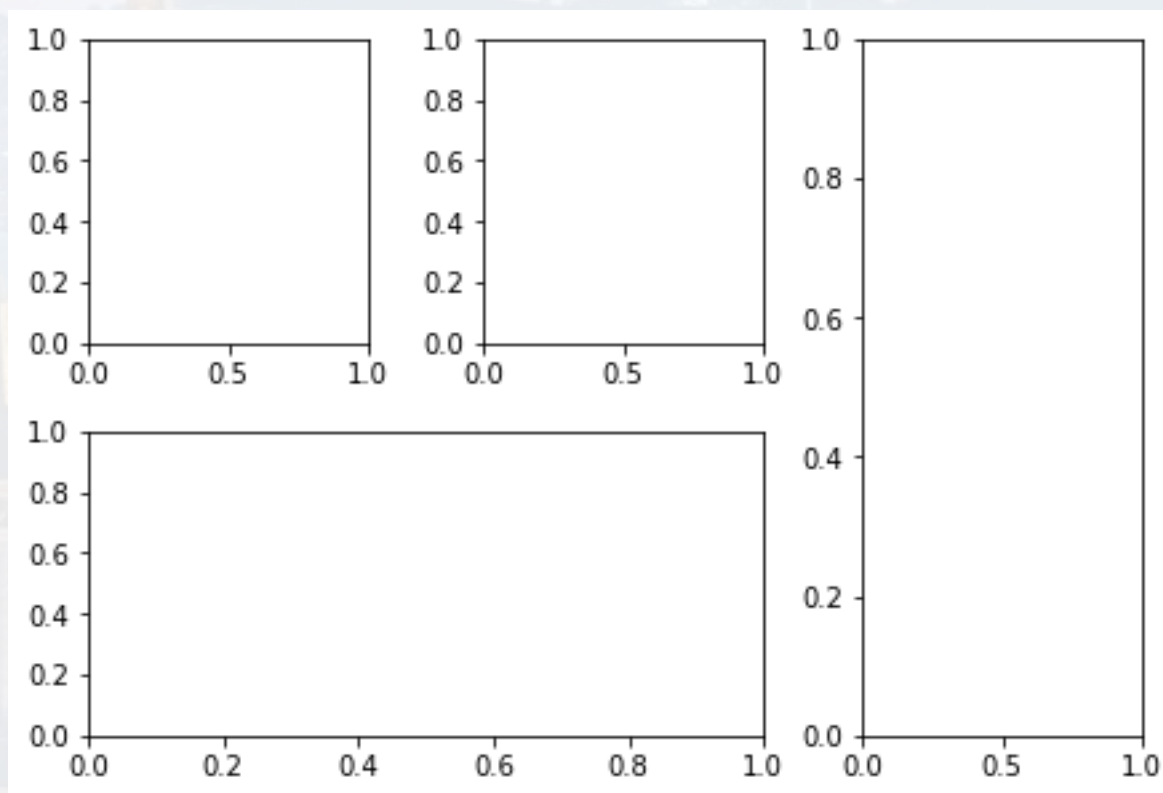
fig

referring to a figure (which can include numerous subplots)

mosaic subplots

classical subplots

```
figMo, axes = plt.subplot_mosaic([[ 'A', 'B', 'C'],\n                                   [ 'D', 'D', 'C']], layout = "constrained")
```





Three levels in Python:

plt

the plot itself

ax

referring to axis of specific plot

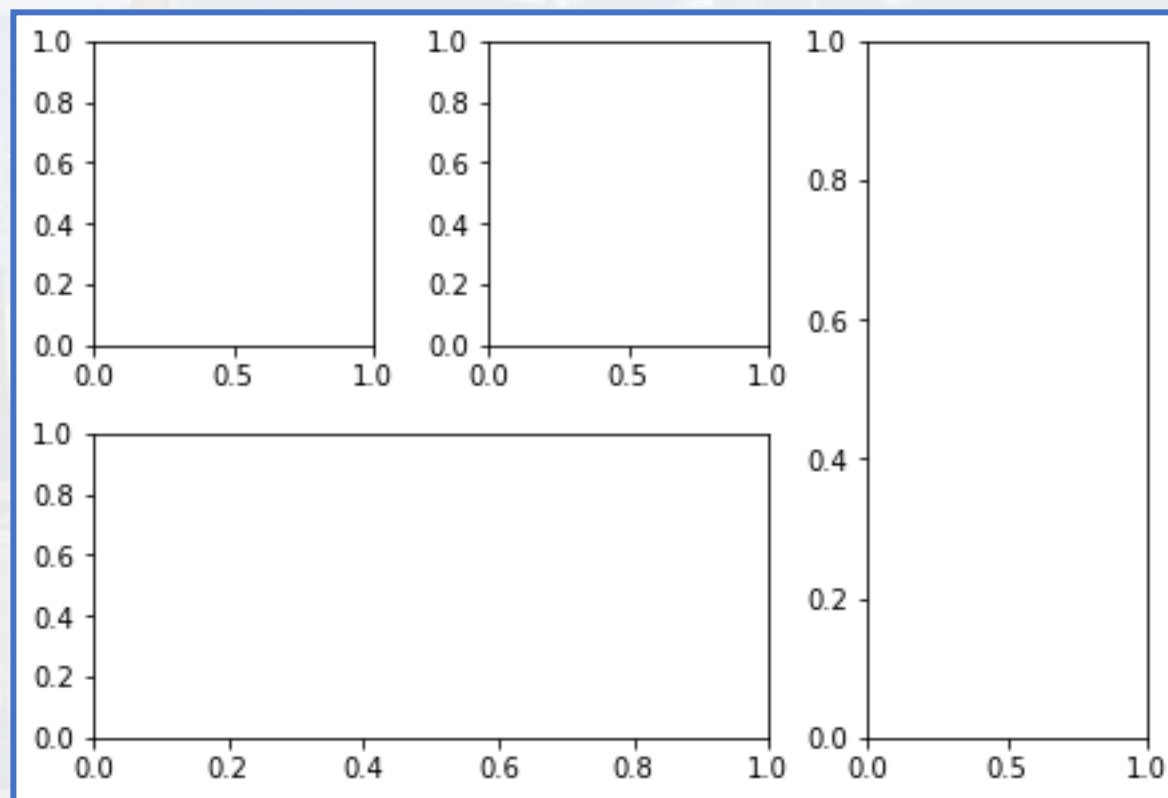
fig

referring to a figure (which can include numerous subplots)

mosaic subplots

classical subplots

```
figMo, axes = plt.subplot_mosaic([['A', 'B', 'C'],\n                                   ['D', 'D', 'C']], layout = "constrained")
```



figMo,

points to the current
figure

Three levels in Python:

plt

the plot itself

ax

referring to axis of specific plot

fig

referring to a figure

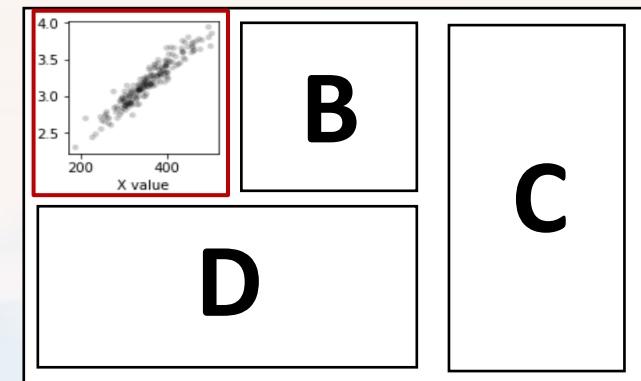
mosaic subplots

classical subplots

```
figMo, axes = plt.subplot_mosaic([[ 'A', 'B', 'C'],\
                                   [ 'D', 'D', 'C']], layout = "constrained")
```

```
axes[ 'A'].scatter(x, y, s = 20, c = 'k', alpha = 0.2, edgecolors = 'none')
axes[ 'A'].set(xlabel = 'X value')
```

populating the
first axis object
with a **scatter plot**





Three levels in Python:

plt

the plot itself

ax

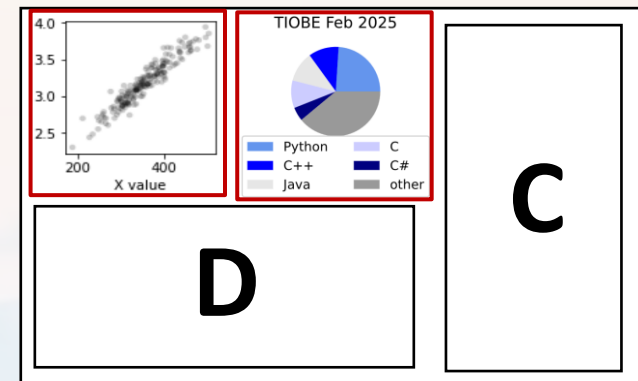
referring to axis of specific plot

fig

referring to a figure

mosaic subplots

classical subplots



```
figMo, axes = plt.subplot_mosaic([[ 'A', 'B', 'C'],\
                                   [ 'D', 'D', 'C']], layout = "constrained")
```

```
axes[ 'A'].scatter(x, y, s = 20, c = 'k', alpha = 0.2, edgecolors = 'none')
axes[ 'A'].set(xlabel = 'X value')
```

```
axes[ 'B'].pie([24, 11, 11, 10, 5, 39],\
               colors = [ '#6495ED', 'blue', [0.9, 0.9, 0.9], '#CCCCFF',\
                          '#000080', '#999999'])
axes[ 'B'].set(title = 'TIOBE Feb 2025')
axes[ 'B'].legend([ 'Python', 'C++', 'Java', 'C', 'C#', 'other'],\
                  bbox_to_anchor = (0.5,-0.5), loc = 'lower center',\
                  ncol = 2)
```

populating the
second axis object
with a **pie chart**



Three levels in Python:

plt

the plot itself

ax

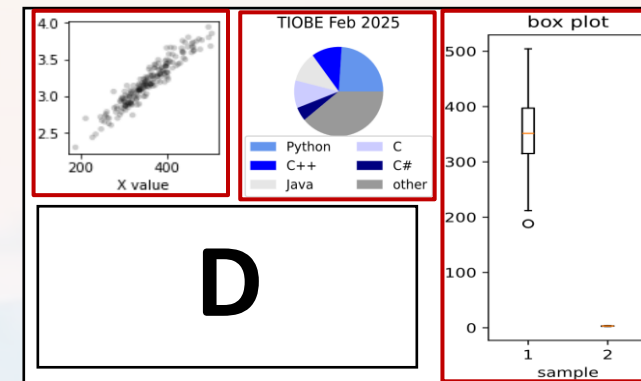
referring to axis of specific plot

fig

referring to a figure

mosaic subplots

classical subplots



```
figMo, axes = plt.subplot_mosaic([[ 'A', 'B', 'C'],\
                                   [ 'D', 'D', 'C']], layout = "constrained")
```

```
axes[ 'A'].scatter(x, y, s = 20, c = 'k', alpha = 0.2, edgecolors = 'none')
axes[ 'A'].set(xlabel = 'X value')
```

```
axes[ 'B'].pie([24, 11, 11, 10, 5, 39], colors = [ '#6495ED', 'blue', [0.9, 0.9, 0.9], '#CCCCFF',\
                                                  '#000080', '#999999'])
```

```
axes[ 'B'].set(title = 'TIOBE Feb 2025')
axes[ 'B'].legend([ 'Python', 'C++', 'Java', 'C', 'C#', 'other'],\
                  bbox_to_anchor = (0.5,-0.5), loc = 'lower center', ncol = 2)
```

and so on...

```
axes[ 'C'].boxplot([x,y])
axes[ 'C'].set(xlabel = 'sample')
axes[ 'C'].set(ylabel = 'values')
axes[ 'C'].set(title = 'box plot')
```



Three levels in Python:

plt

the plot itself

ax

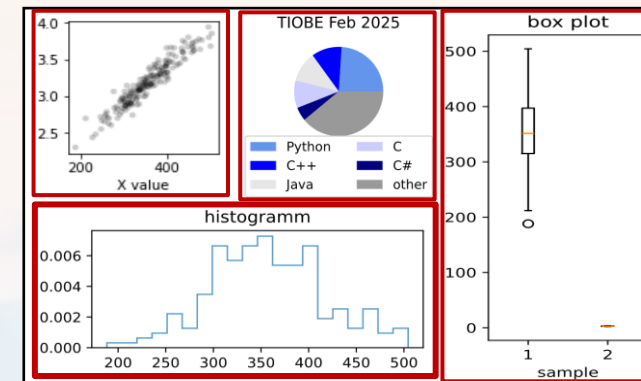
referring to axis of specific plot

fig

referring to a figure

mosaic subplots

classical subplots



```
figMo, axes = plt.subplot_mosaic([[ 'A', 'B', 'C'],\
                                   [ 'D', 'D', 'C']], layout = "constrained")
```

```
axes[ 'A'].scatter(x, y, s = 20, c = 'k', alpha = 0.2, edgecolors = 'none')
axes[ 'A'].set(xlabel = 'X value')
```

```
axes[ 'B'].pie([24, 11, 11, 10, 5, 39], colors = [ '#6495ED', 'blue', [0.9, 0.9, 0.9], '#CCCCFF',\
                                                  '#000080', '#999999'])
```

```
axes[ 'B'].set(title = 'TIOBE Feb 2025')
axes[ 'B'].legend([ 'Python', 'C++', 'Java', 'C', 'C#', 'other'],\
                  bbox_to_anchor = (0.5,-0.5), loc = 'lower center', ncol = 2)
```

```
axes[ 'C'].boxplot([x,y])
axes[ 'C'].set(xlabel = 'sample')
axes[ 'C'].set(ylabel = 'values')
axes[ 'C'].set(title = 'box plot')
```

```
axes[ 'D'].hist(x, 20, density = True, histtype = 'step', facecolor = 'g',\
               alpha = 0.75)
```

```
axes[ 'D'].set(title = 'histogram')
```

and so on...



Three levels in Python:

plt

the plot itself

ax

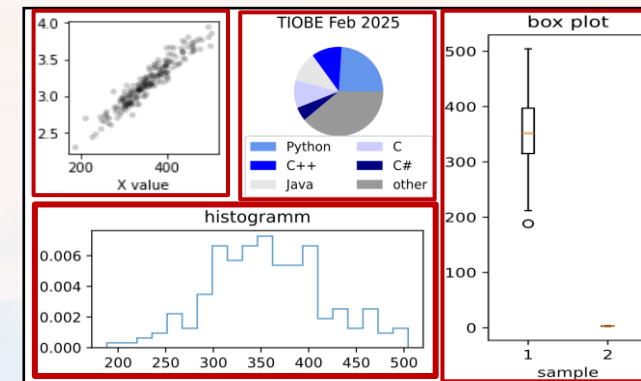
referring to axis of specific plot

fig

referring to a figure

mosaic subplots

classical subplots



reference to the
individual axes

```
figMo, axes = plt.subplot_mosaic([[ 'A', 'B', 'C'],\
                                   [ 'D', 'D', 'C']], layout = "constrained")
```

```
axes[ 'A'].scatter(x, y, s = 20, c = 'k', alpha = 0.2, edgecolors = 'none')
axes[ 'A'].set(xlabel = 'X value')
```

```
axes[ 'B'].pie([24, 11, 11, 10, 5, 39], colors = [ '#6495ED', 'blue', [0.9, 0.9, 0.9], '#CCCCFF',\
                                                  '#000080', '#999999'])
```

```
axes[ 'B'].set(title = 'TIOBE Feb 2025')
```

```
axes[ 'B'].legend([ 'Python', 'C++', 'Java', 'C', 'C#', 'other'],\
                  bbox_to_anchor = (0.5,-0.5), loc = 'lower center', ncol = 2)
```

```
axes[ 'C'].boxplot([x,y])
```

```
axes[ 'C'].set(xlabel = 'sample')
```

```
axes[ 'C'].set(ylabel = 'values')
```

```
axes[ 'C'].set(title = 'box plot')
```

```
axes[ 'D'].hist(x, 20, density = True, histtype = 'step', facecolor = 'g',\
               alpha = 0.75)
```

```
axes[ 'D'].set(title = 'histogram')
```




Three levels in Python:

plt

the plot itself

ax

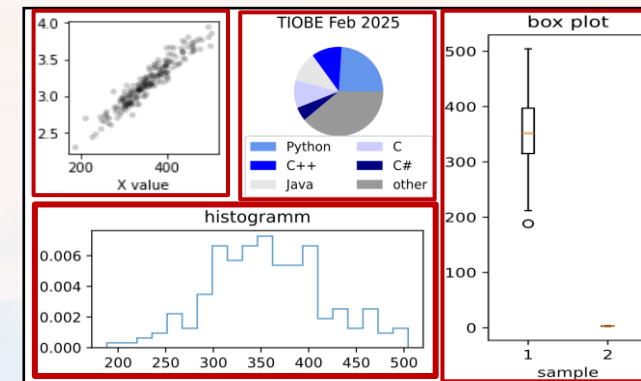
referring to axis of specific plot

fig

referring to a figure

mosaic subplots

classical subplots



```
figMo, axes = plt.subplot_mosaic([[ 'A', 'B', 'C'],\
                                   [ 'D', 'D', 'C']], layout = "constrained")
```

```
axes[ 'A'].scatter(x, y, s = 20, c = 'k', alpha = 0.2, edgecolors = 'none')
axes[ 'A'].set(xlabel = 'X value')
```

```
axes[ 'B'].pie([24, 11, 11, 10, 5, 39], colors = [ '#6495ED', 'blue', [0.9, 0.9, 0.9], '#CCCCFF',\
                                                  '#000080', '#999999'])
```

```
axes[ 'B'].set(title = 'TIOBE Feb 2025')
axes[ 'B'].legend([ 'Python', 'C++', 'Java', 'C', 'C#', 'other'],\
                  bbox_to_anchor = (0.5,-0.5), loc = 'lower center', ncol = 2)
```

```
axes[ 'C'].boxplot([x,y])
axes[ 'C'].set(xlabel = 'sample')
axes[ 'C'].set(ylabel = 'values')
axes[ 'C'].set(title = 'box plot')
```

```
axes[ 'D'].hist(x, 20, density = True, histtype = 'step', facecolor = 'g',\
               alpha = 0.75)
axes[ 'D'].set(title = 'histogram')
```

notice the
different color
codes



Three levels in Python:

plt

the plot itself

ax

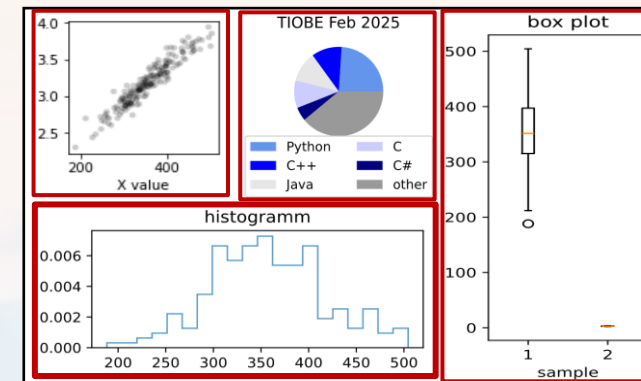
referring to axis of specific plot

fig

referring to a figure

mosaic subplots

classical subplots



```
figMo, axes = plt.subplot_mosaic([[ 'A', 'B', 'C'],\
                                   [ 'D', 'D', 'C']], layout = "constrained")
```

```
axes[ 'A'].scatter(x, y, s = 20, c = 'k', alpha = 0.2, edgecolors = 'none')
axes[ 'A'].set(xlabel = 'X value')
```

...

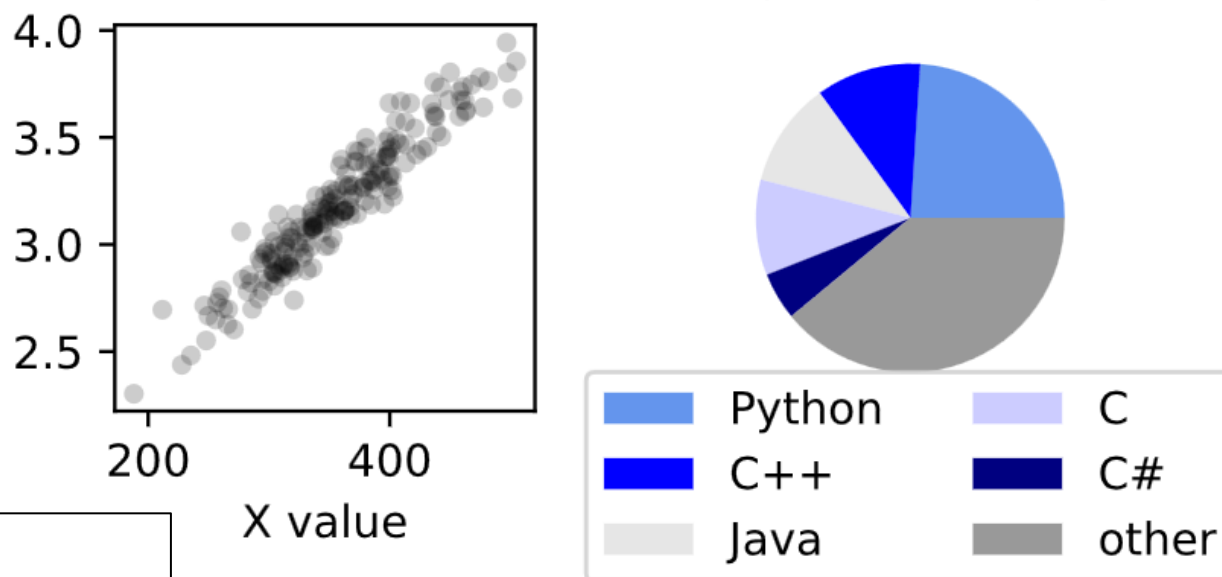
```
axes[ 'D'].hist(x, 20, density = True, histtype = 'step', facecolor = 'g',\
               alpha = 0.75)
axes[ 'D'].set(title = 'histogram')
```

```
figMo.savefig('test.pdf', dpi = 1600)
```

referring to the
specific figure

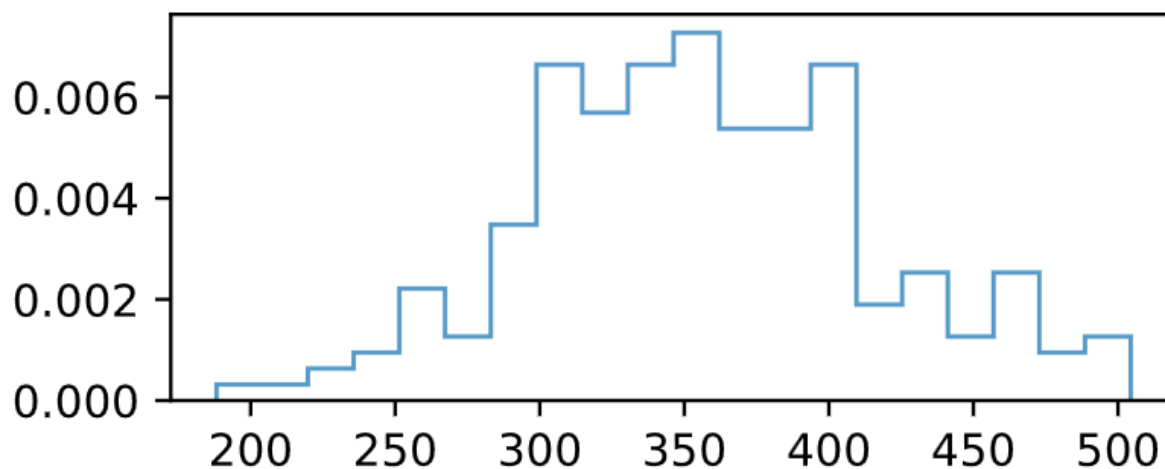


TIOBE Feb 2025

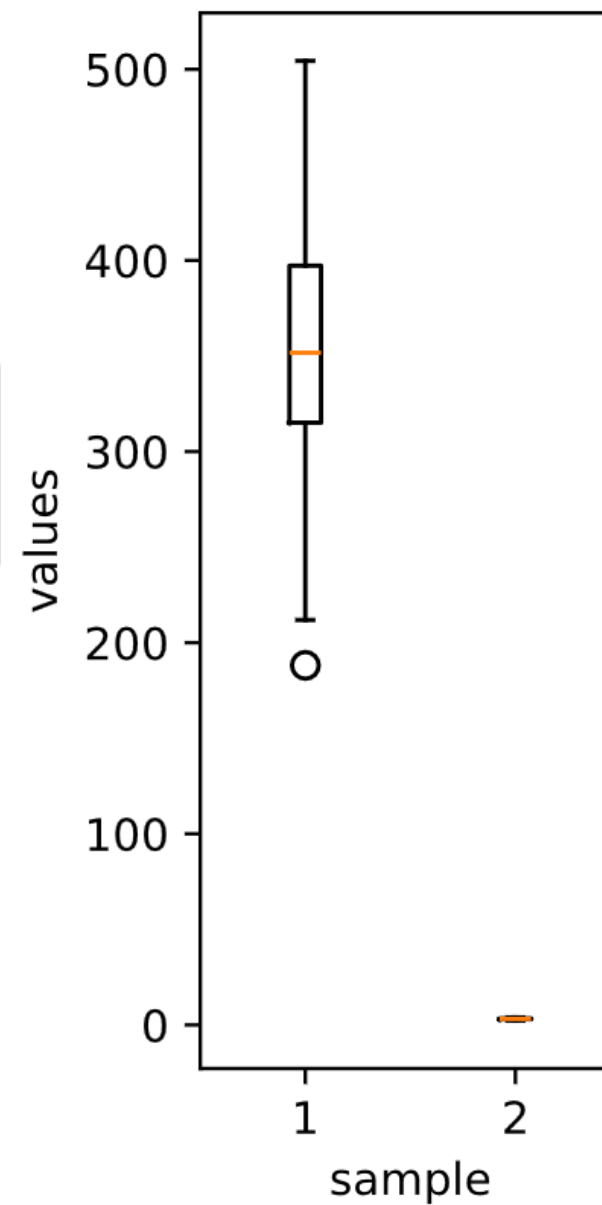


check out
`PlotMosaicExample.py`

histogramm



box plot





Three levels in Python:

plt

the plot itself

ax

referring to axis of specific plot

fig

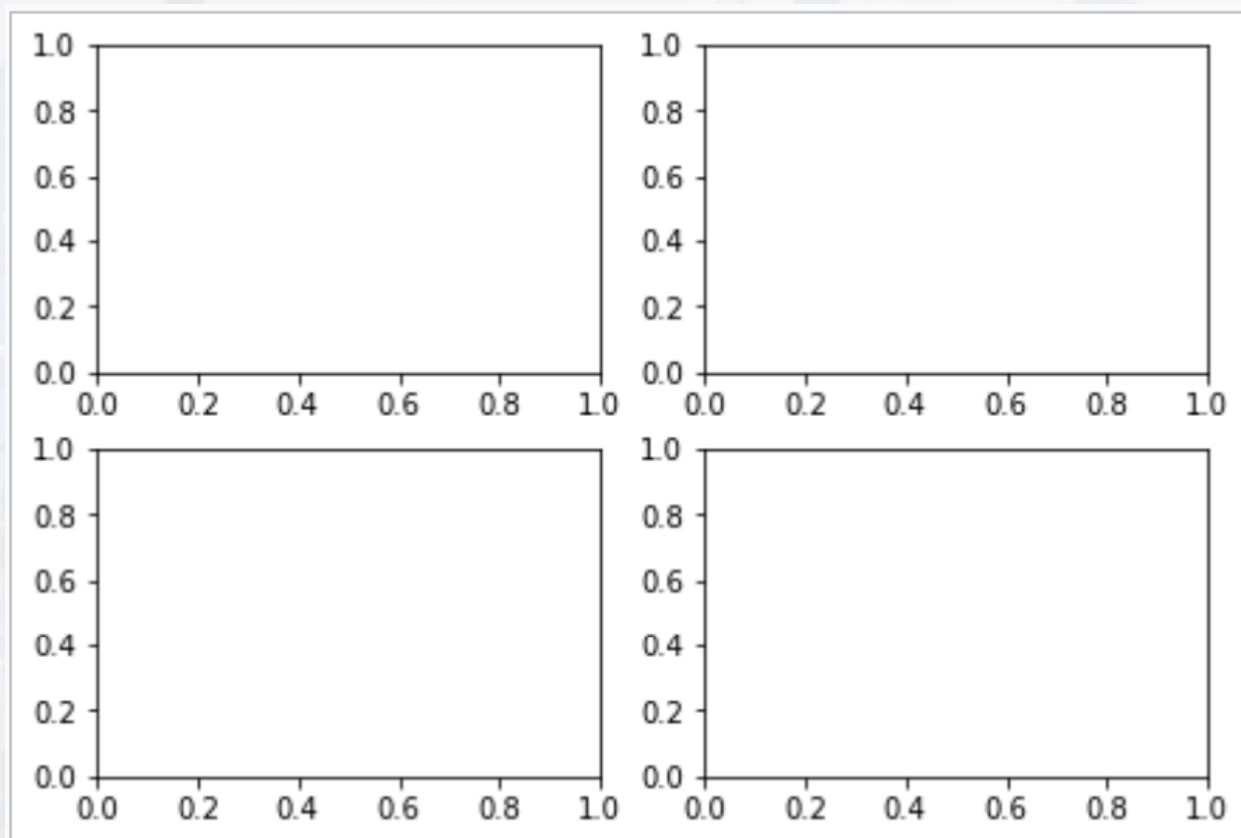
referring to a figure (which can include numerous subplots)

mosaic subplots

classical subplots

Same idea!

```
figSt, axes = plt.subplots(nrows = 2, ncols = 2, layout = "constrained")
```





Three levels in Python:

plt

the plot itself

ax

referring to axis of specific plot

fig

referring to a figure (which can include numerous subplots)

mosaic subplots

classical subplots

Same idea!

in mosaic mode:
label can be
numeric or **str**.
here: has to be
int (refers to
location in
figure)

```
figSt, axes = plt.subplots(nrows = 2, ncols = 2, layout = "constrained")
```

```
axes[0, 0].scatter(x, y, s = 20, c = 'k', alpha = 0.2, edgecolors = 'none')
```

```
axes[0, 0].set(xlabel = 'X value')
```

```
axes[1, 0].pie([24, 11, 11, 10, 5, 39], colors = ['#6495ED', 'blue', [0.9, 0.9, 0.9],\n          '#CCCCFF', '#000080', '#999999'])
```

```
axes[1, 0].set(title = 'TIOBE Feb 2025')
```

```
axes[1, 0].legend(['Python', 'C++', 'Java', 'C', 'C#', 'other'],\n                  bbox_to_anchor = (0.5, -0.5), loc = 'lower center', ncol = 2)
```

```
axes[0, 1].boxplot([x, y])
```

```
axes[0, 1].set(xlabel = 'sample')
```

```
axes[0, 1].set(ylabel = 'values')
```

```
axes[0, 1].set(title = 'box plot')
```

```
axes[1, 1].hist(x, 20, density = True, histtype = 'step', facecolor = 'g', alpha = 0.75)
```

```
axes[1, 1].set(title = 'histogram')
```



Three levels in Python:

plt

the plot itself

ax

referring to axis of specific plot

fig

referring to a figure (which can include numerous subplots)

mosaic subplots

classical subplots

Same idea!

```
figSt, axes = plt.subplots(nrows = 2, ncols = 2, layout = "constrained")
```

```
axes[0, 0].scatter(x, y, s = 20, c = 'k', alpha = 0.2, edgecolors = 'none')
```

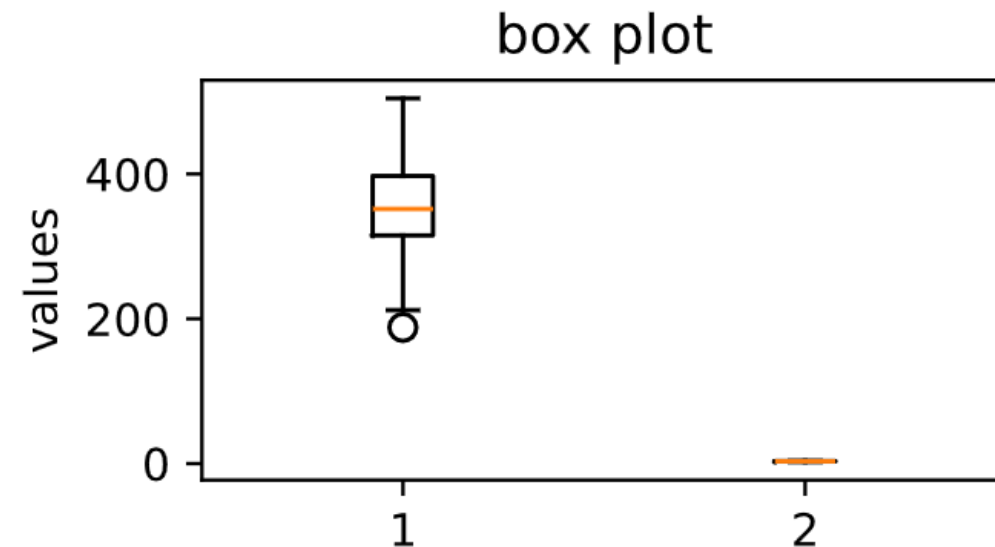
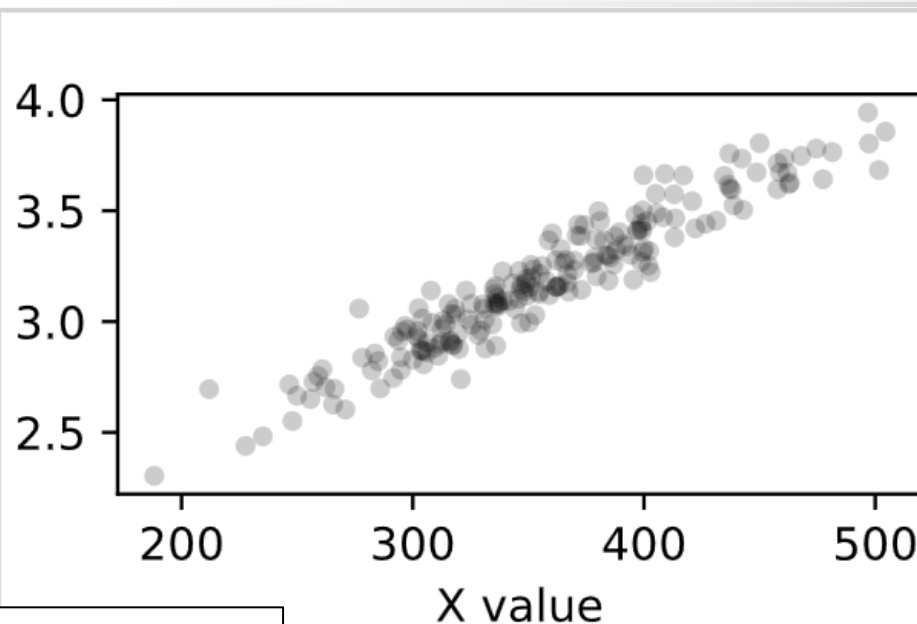
```
axes[0, 0].set(xlabel = 'X value')
```

...

```
axes[1, 1].hist(x, 20, density=True, histtype = 'step', facecolor = 'g', alpha = 0.75)
```

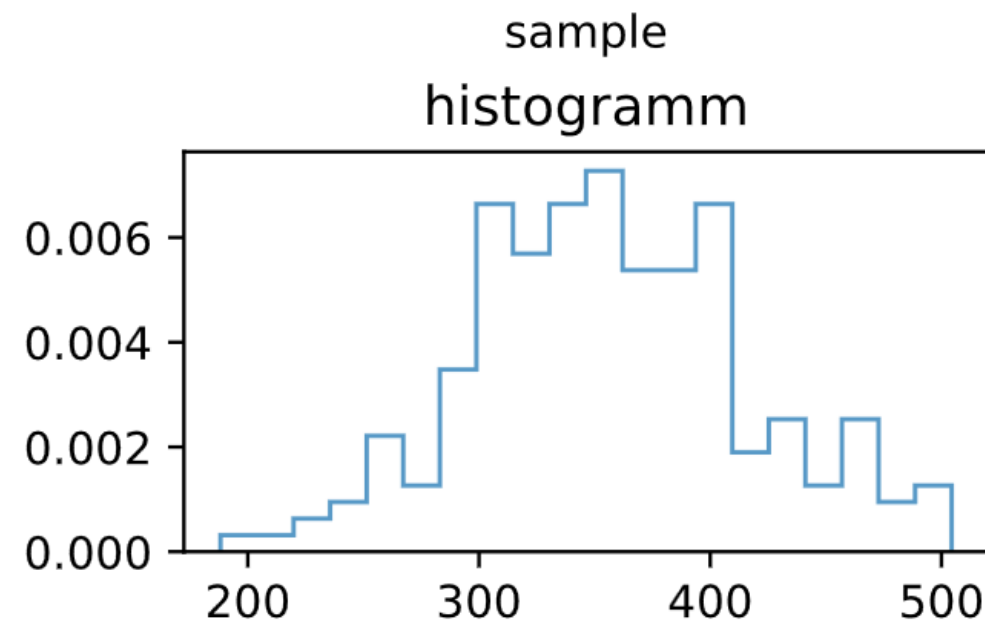
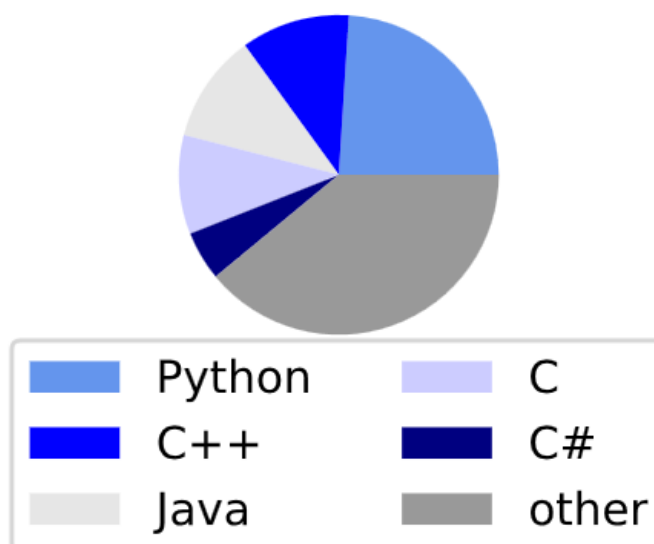
```
axes[1, 1].set(title = 'histogramm')
```

```
figSt.savefig('test.pdf', dpi = 1600)
```

check out
`PlotStandardExample.py`

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Thank you for your attention!

