Plotting and Visualization

Using data visualization libraries

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matplotlib API primer

- Figures and subplots
- adjusting the spacing around subplots
- colors, markers and line styles
- ticks, labels, and legends
- adding legends
- annotation and drawing on a subplot
- saving plots to file
- matplotlib configuration

plotting with pandas and seaborn

• line plots

- Bar plots
- Histogram and density plots
- Scatter or Point plots
- Facet Grids and Categorical Data

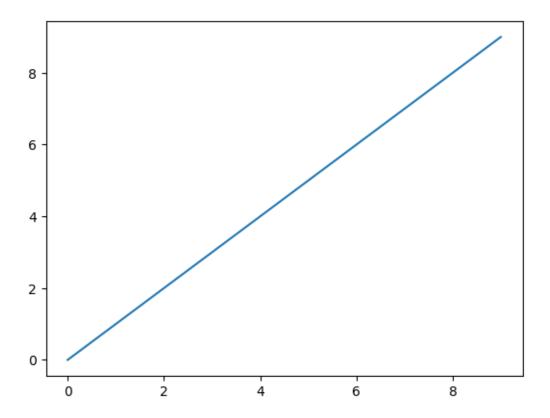
Into matplotlib

```
import matplotlib.pyplot as plt
import numpy as np
import pandas as pd
%matplotlib inline

data = np.arange(10)
data

array([0, 1, 2, 3, 4, 5, 6, 7, 8, 9])

plt.plot(data)
```



Figures and subplots

```
fig = plt.figure()
plt.show()

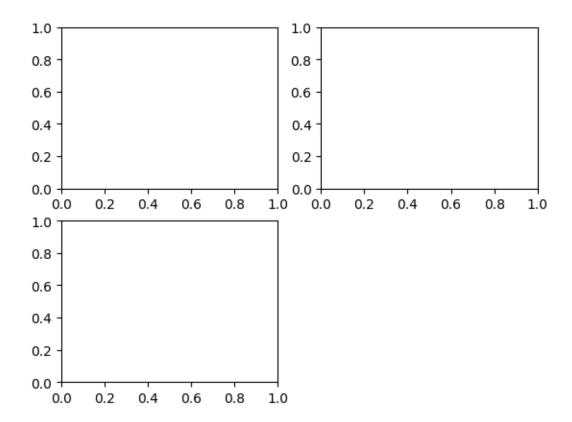
# 2, 2 means 4 sub-plots will be created

# % matplotlib
ax1 = fig.add_subplot(2, 2, 1)

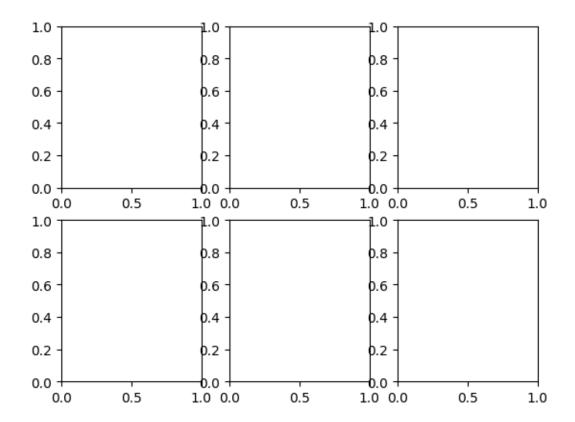
ax2 = fig.add_subplot(2, 2, 2)
ax3 = fig.add_subplot(2, 2, 3)

fig = plt.figure()
ax1 = fig.add_subplot(2, 2, 1)
```

```
ax2 = fig.add_subplot(2, 2, 2)
ax3 = fig.add_subplot(2, 2, 3)
```

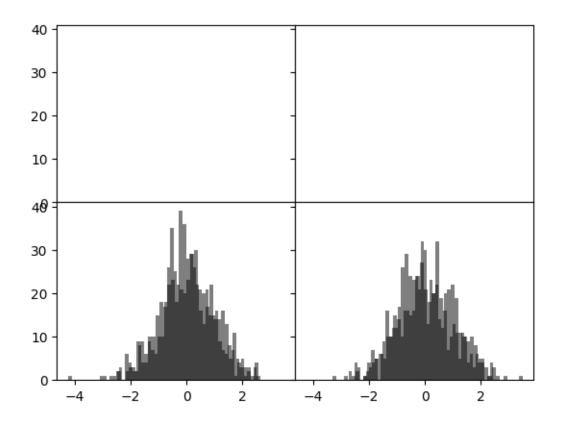


```
! pip install ipympl
fig, axes = plt.subplots(2,3)
```

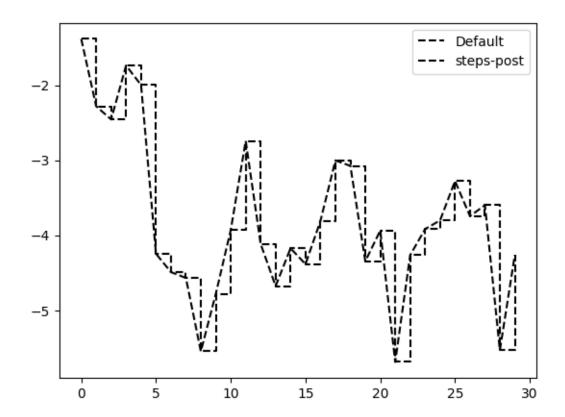


axes

Adjusting spacing around subplots



<matplotlib.legend.Legend at 0x205861b2460>

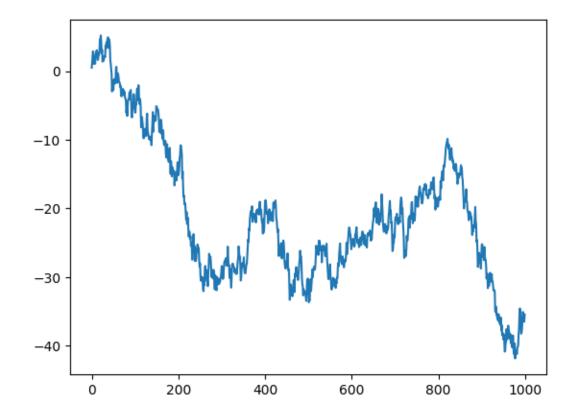


Ticks, Labels, and Legends

```
ax.get_xlim([0, 10])
  help(plt.xlim)
Help on function xlim in module matplotlib.pyplot:
xlim(*args, **kwargs)
    Get or set the x limits of the current axes.
    Call signatures::
        left, right = xlim() # return the current xlim
        xlim((left, right)) # set the xlim to left, right
        xlim(left, right) # set the xlim to left, right
    If you do not specify args, you can pass *left* or *right* as kwargs,
    i.e.::
        xlim(right=3) # adjust the right leaving left unchanged
        xlim(left=1) # adjust the left leaving right unchanged
    Setting limits turns autoscaling off for the x-axis.
    Returns
    left, right
        A tuple of the new x-axis limits.
    Notes
    Calling this function with no arguments (e.g. ``xlim()``) is the pyplot
    equivalent of calling `~.Axes.get_xlim` on the current axes.
    Calling this function with arguments is the pyplot equivalent of calling
    `~.Axes.set_xlim` on the current axes. All arguments are passed though.
```

Setting the title, axis labels, ticks, and tick labels

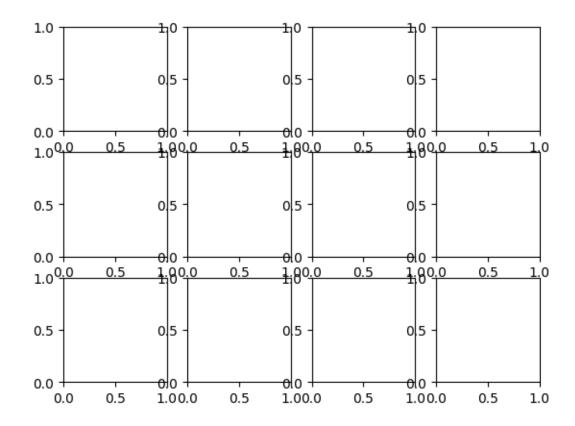
```
fix, ax = plt.subplots()
ax.plot(np.random.standard_normal(1000).cumsum());
```



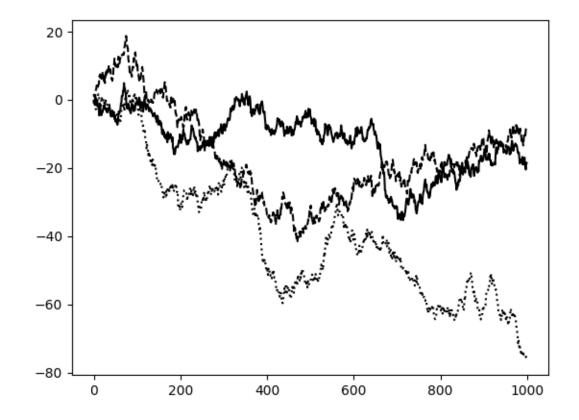
```
Text(0.5, 1.0, 'My matplotlib plot')
   ax.set_title('My matplotlib plot')
Text(0.5, 1.0, 'My matplotlib plot')
   plt.show()
```

Adding legends

```
fig, ax = plt.subplots(3, 4)
```



```
fig,ax = plt.subplots()
ax.plot(np.random.randn(1000).cumsum(), color = 'black',
```

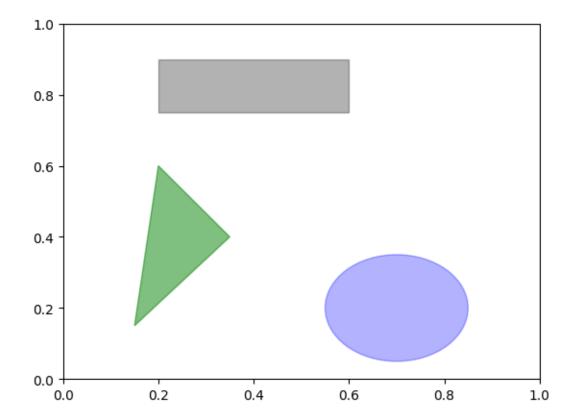


ax.legend()

<matplotlib.legend.Legend at 0x20588a6e700>

Annotations and Drawing on a Subplot

<matplotlib.patches.Polygon at 0x20588b71a60>



Saving photos to file

```
fig.savefig('figpath.svg')
fig.savefig('figpath.png', dpi=400)
```

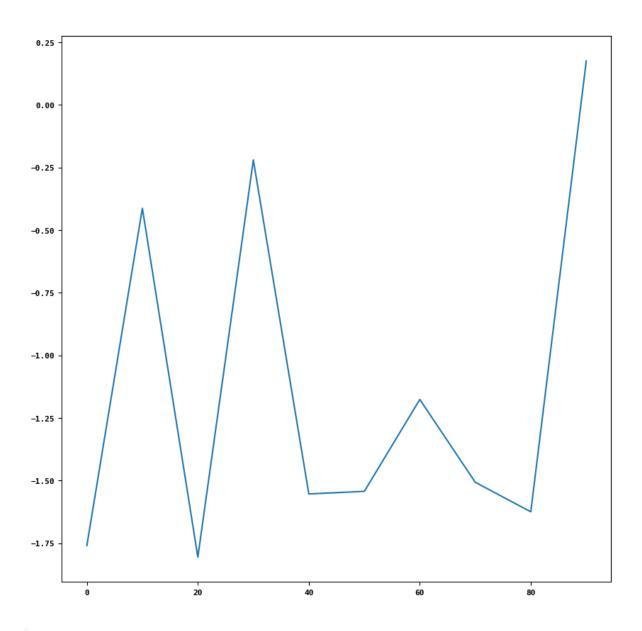
Matplotlib configuration

```
plt.rc('figure', figsize = (10, 10))

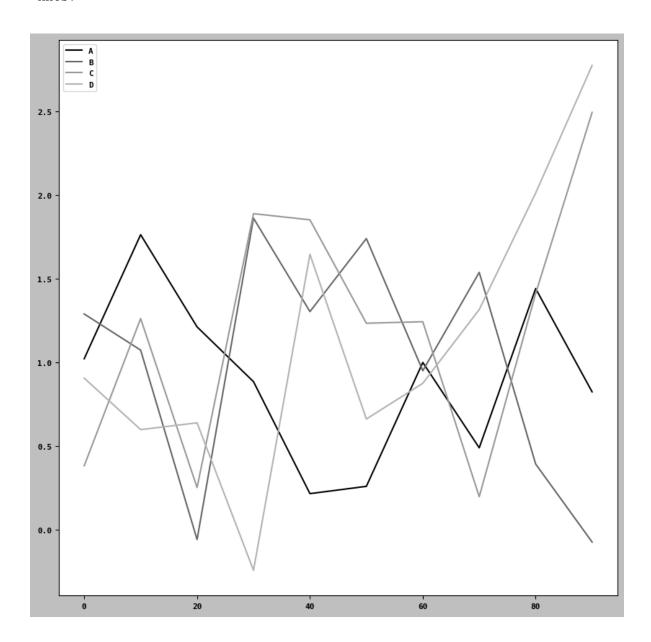
plt.rc('font', family= 'monospace', weight = 'bold', size= 8)
```

Plotting with pandas and seaborn

Line plots



<Axes: >



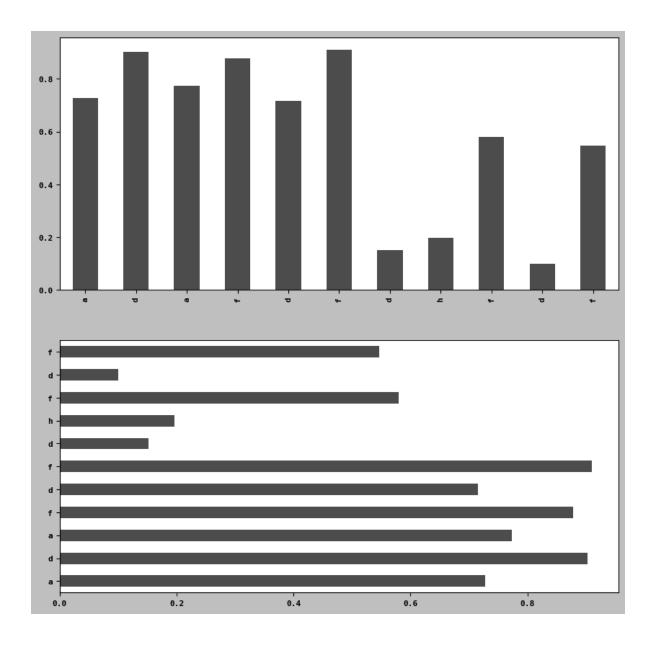
Bar Plots

```
fig, axes = plt.subplots(2, 1)

data = pd.Series(np.random.uniform(size=11),
```

```
index=list('adafdfdhfdf'))
data.plot.bar(ax = axes[0], color= 'black', alpha=0.7)
data.plot.barh(ax= axes[1], color= 'black', alpha= 0.7)
```

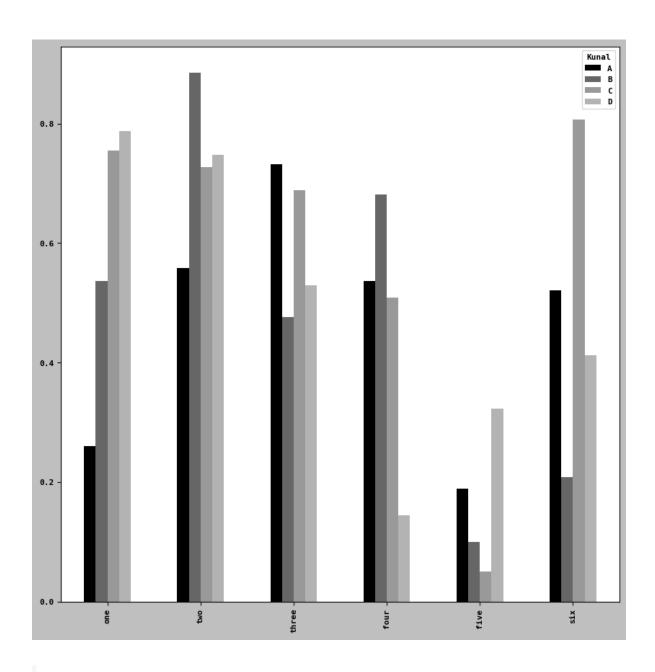
<Axes: >



Kunal	A	В	С	D
one	0.260658	0.536198	0.754708	0.788162
two	0.558069	0.885258	0.726874	0.747412
$_{\mathrm{three}}$	0.731644	0.476325	0.688620	0.529862
four	0.536672	0.681522	0.509112	0.143861
five	0.188829	0.099173	0.050697	0.323006
six	0.521520	0.208836	0.807558	0.411876

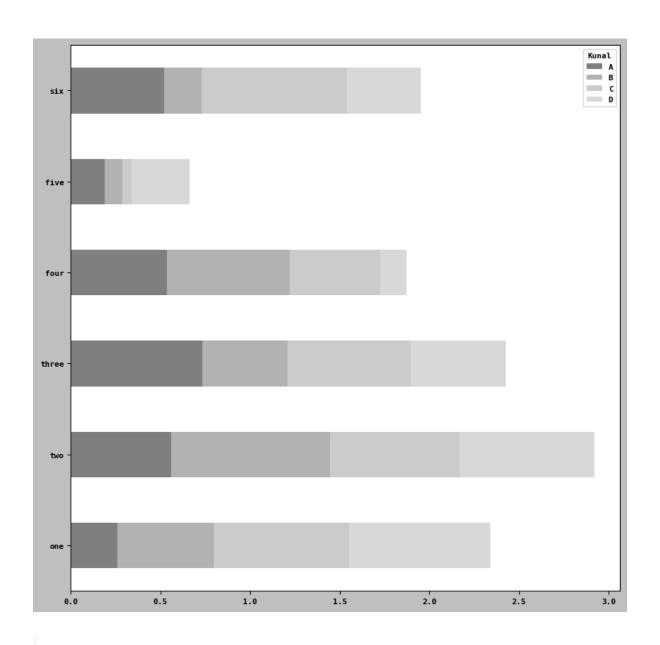
df2.plot.bar()

<Axes: >



df2.plot.barh(stacked=True, alpha=0.5)

<Axes: >



iris = pd.read_csv(r"E:/pythonfordatanalysis/semainedu26fevrier/iris.csv")
iris.head()

	Id	Sepal Length (cm)	Sepal Width (cm)	Petal Length (cm)	Petal Width (cm)	Species
0	1	5.1	3.5	1.4	0.2	Iris-setosa
1	2	4.9	3.0	1.4	0.2	Iris-setosa
2	3	4.7	3.2	1.3	0.2	Iris-setosa
3	4	4.6	3.1	1.5	0.2	Iris-setosa

	Id	Sepal Length (cm)	Sepal Width (cm)	Petal Length (cm)	Petal Width (cm)	Species
$\overline{4}$	5	5.0	3.6	1.4	0.2	Iris-setosa

```
iris.tail()
#len_wd = pd.
```

	Id	Sepal Length (cm)	Sepal Width (cm)	Petal Length (cm)	Petal Width (cm)	Species
145	146	6.7	3.0	5.2	2.3	Iris-virginica
146	147	6.3	2.5	5.0	1.9	Iris-virginica
147	148	6.5	3.0	5.2	2.0	Iris-virginica
148	149	6.2	3.4	5.4	2.3	Iris-virginica
149	150	5.9	3.0	5.1	1.8	Iris-virginica

```
print(iris.columns)
```

```
count = pd.crosstab(iris['Sepal Length (cm)'], iris['Sepal Width (cm)'])
count2 = count.reindex(index=['length', 'width', ])
```

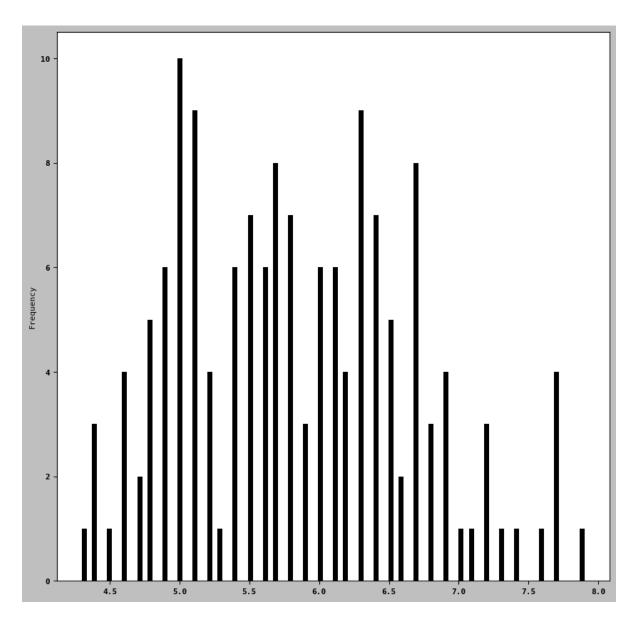
 ${\tt count2}$

Sepal Width (cm)	2.0	2.2	2.3	2.4	2.5	2.6	2.7	2.8	2.9	3.0		3.4	3.5	
Sepal Length (cm)														
length	NaN		NaN	NaN										
width	NaN	•••	NaN	NaN										
														_

Histogram and Density plots

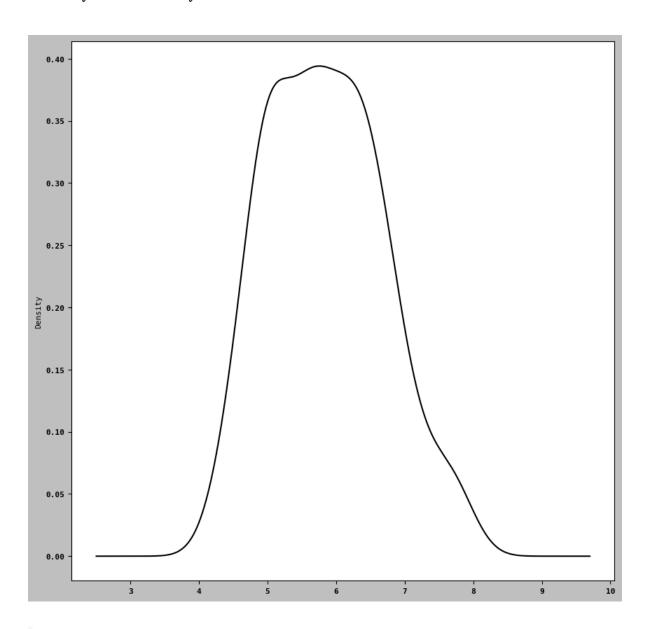
```
iris['Sepal Length (cm)'].plot.hist(bins= 100)
```

<Axes: ylabel='Frequency'>



iris['Sepal Length (cm)'].plot.density()

<Axes: ylabel='Density'>



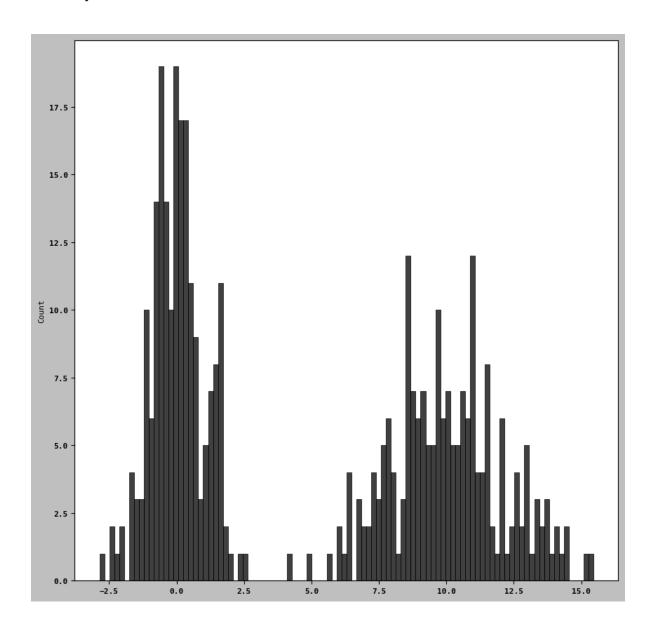
! pip install seaborn

import seaborn as sns

```
comp1 = np.random.standard_normal(200)
comp2 = 10 + 2 * np.random.standard_normal(200)

values = pd.Series(np.concatenate([comp1, comp2]))
sns.histplot(values, bins= 100, color = 'black')
```

<Axes: ylabel='Count'>



Scatter or point plots

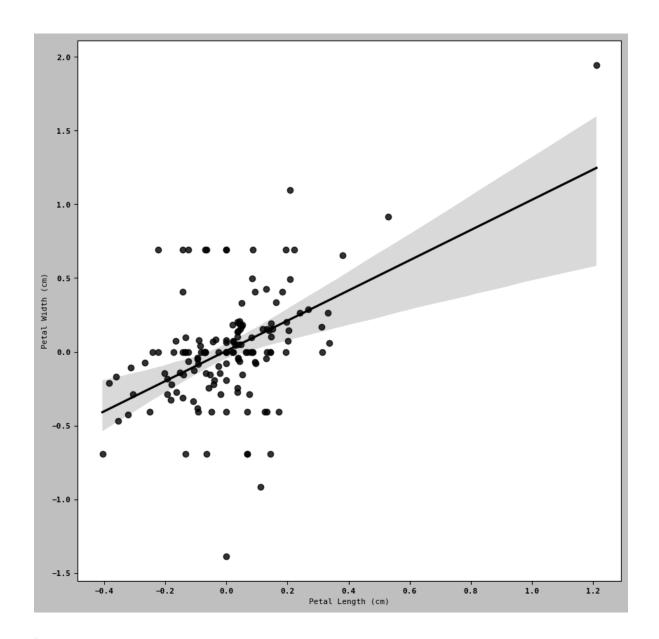
iris.head()

	Id	Sepal Length (cm)	Sepal Width (cm)	Petal Length (cm)	Petal Width (cm)	Species
0	1	5.1	3.5	1.4	0.2	Iris-setosa
1	2	4.9	3.0	1.4	0.2	Iris-setosa
2	3	4.7	3.2	1.3	0.2	Iris-setosa
3	4	4.6	3.1	1.5	0.2	Iris-setosa
4	5	5.0	3.6	1.4	0.2	Iris-setosa

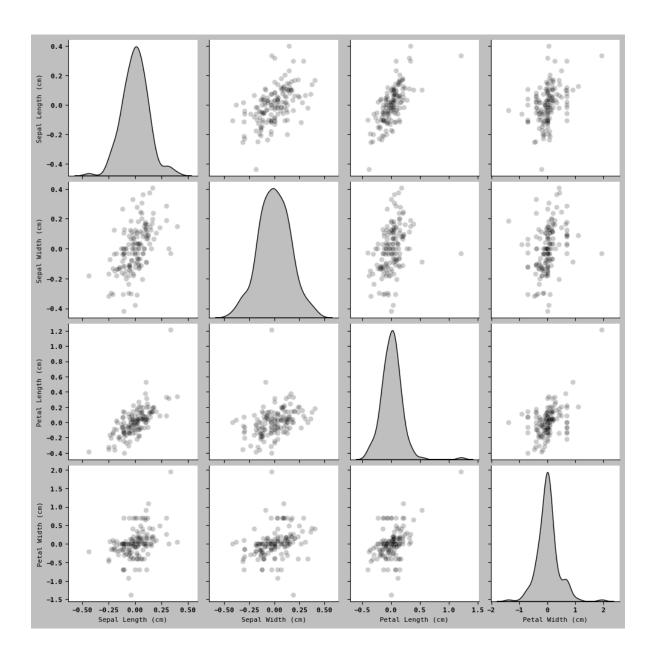
	Sepal Length (cm)	Sepal Width (cm)	Petal Length (cm)	Petal Width (cm)
145	0.000000	-0.095310	-0.091808	-0.083382
146	-0.061558	-0.182322	-0.039221	-0.191055
147	0.031253	0.182322	0.039221	0.051293
148	-0.047253	0.125163	0.037740	0.139762
149	-0.049597	-0.125163	-0.057158	-0.245122

using regplot method to make scatter plots

```
ax = sns.regplot(x= "Petal Length (cm)", y = "Petal Width (cm)", data= trans_iris2)
#ax.title("Change in log (Petal Length (cm)) length versus log (Petal Width (cm)) width ")
```



 $sns.pairplot(trans_iris2, \ diag_kind= \ 'kde', \ plot_kws=\{'alpha': \ {\tt 0.2}\}\)$



Facet Grids and Categorical Data

• catplots

SyntaxError: invalid syntax (3918704872.py, line 2)

```
# renaming columns

df3 = df.rename(columns={
    'Sepal Length (cm)': 'sepal_lengh',
    'Sepal Width (cm)': 'sepal_width_1',
    'Petal Length (cm)': 'petal_length',
    'Petal Width (cm)': 'petal_width_2',
})

df3
```

	A	В	\mathbf{C}	D
0	1.023099	1.290412	0.383457	0.906869
10	1.764172	1.074479	1.263072	0.599487
20	1.213259	-0.057754	0.253086	0.639868
30	0.885836	1.862965	1.889980	-0.241599
40	0.216726	1.304783	1.853073	1.646926
50	0.260248	1.741218	1.235233	0.662542
60	1.000675	0.951141	1.243977	0.876234
70	0.490475	1.539086	0.198194	1.316823
80	1.442176	0.393618	1.413128	2.013297
90	0.825008	-0.072619	2.495192	2.775830

