Matplotlib

- Jake VanderPlas. 2016. Python Data Science Handbook: Essential Tools for Working with Data. O'Reilly Media, Inc.
- Chapter 4 Visualization with Matplotlib
- https://github.com/jakevdp/PythonDataScienceHandbook
- NOTE: many code-examples are deprecated

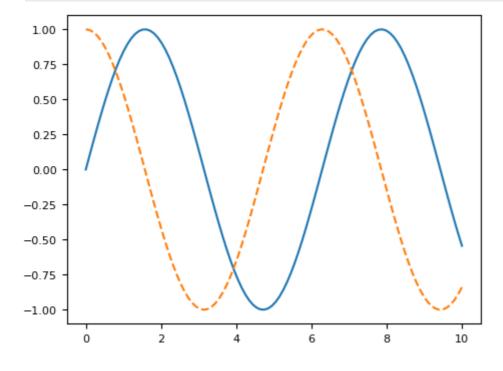
Matplotlib provides:

- Easy and customizable type of plots (line, scatter, bars, histogram, contour, 3D...)
- Integration with NumPy and Pandas
- Extensible rendering using third-party libraries (Seaborn or Plotly)
- Different output formats
 - Static (%matplotlib inline): PNG, JPG, SVG, PDF
 - Interactive (%matplotlib notebook): Qt, Tkinter and WebAgg backends

```
import matplotlib as mpl
import matplotlib.pyplot as plt
import seaborn as sns
import numpy as np
import pandas as pd

# Set the default figure size in inches (width, height)
mpl.rcParams['figure.figsize'] = (5.33,4)
# Set the default font sizes for axes ant tick labels in points
mpl.rcParams['axes.labelsize'] = 10 # Example: 14 points
mpl.rcParams['xtick.labelsize'] = 8 # Example: 12 points for x-axis ticks
mpl.rcParams['ytick.labelsize'] = 8 # Example: 12 points for y-axis ticks
```

```
In [2]: x = np.linspace(0, 10, 100)
fig = plt.figure()
plt.plot(x, np.sin(x), '-')
plt.plot(x, np.cos(x), '--');
```



Dual Interface

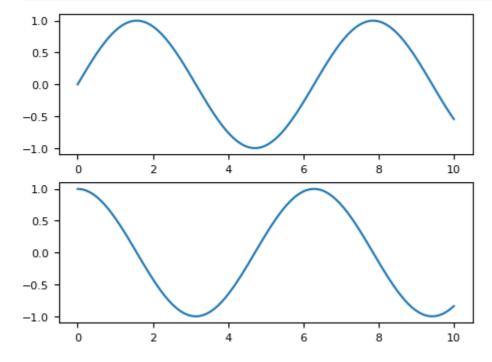
- MATLAB-style Interface
 - Matplotlib: Python alternative for MATLAB
 - Use plt for almost everything
 - Convenient for simple plots
- Object-oriented interface
 - Create objects and call methods on them
 - More flexible

MATLAB-style Interface:

```
In [3]: plt.figure() # create a plot figure

# create the first of two panels and set current axis
plt.subplot(2, 1, 1) # (rows, columns, panel number)
plt.plot(x, np.sin(x))

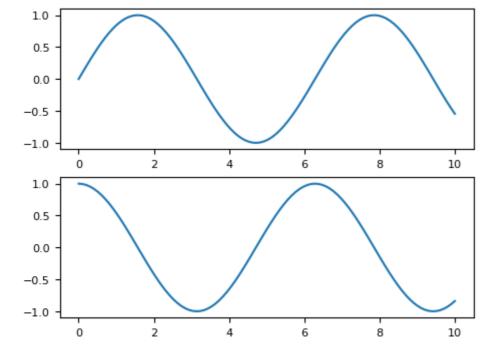
# create the second panel and set current axis
plt.subplot(2, 1, 2)
plt.plot(x, np.cos(x));
```



Object-oriented Interface:

```
In [4]: # First create a grid of plots
# ax will be an array of two Axes objects
fig, ax = plt.subplots(2,1)

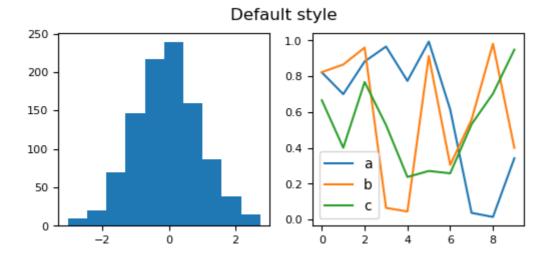
# Call plot() method on the appropriate object
ax[0].plot(x, np.sin(x))
ax[1].plot(x, np.cos(x));
```



Styles

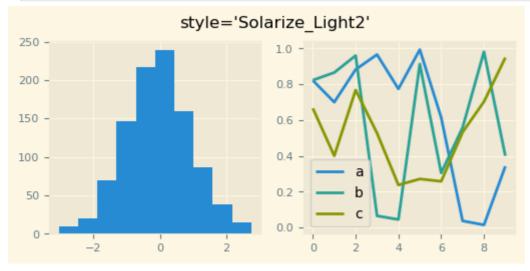
- Predefined collection of visual attributes
- plt.style.library → available styles an their parameters
 - plt.style.available == sorted(plt.style.library.keys())
- plt.style.use('stylename') → set global style
 - Use context manager to chancge the style locallY:
 - with plt.style.context('stylename'):
 make_a_plot()

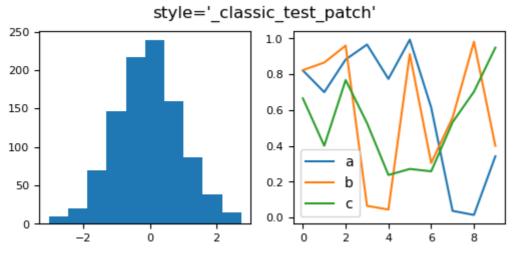
In [6]: hist_and_lines()

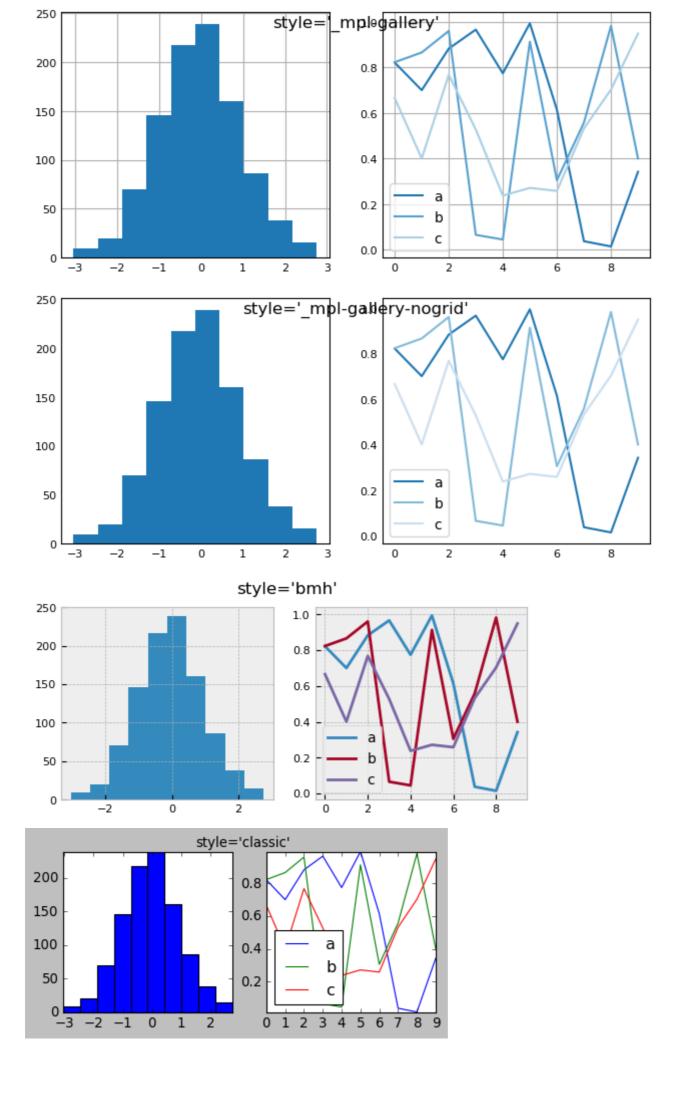


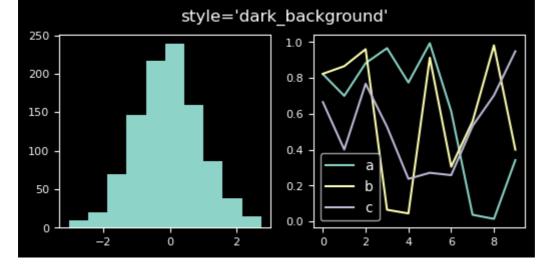
```
Out[7]: ['Solarize_Light2',
           '_classic_test_patch',
          '_mpl-gallery',
          '_mpl-gallery-nogrid',
          'bmh',
          'classic',
          'dark_background',
          'fast',
          'fivethirtyeight',
          'ggplot',
          'grayscale',
          'petroff10',
          'seaborn-v0_8',
          'seaborn-v0_8-bright',
          'seaborn-v0_8-colorblind',
          'seaborn-v0_8-dark',
          'seaborn-v0_8-dark-palette',
          'seaborn-v0_8-darkgrid',
          'seaborn-v0_8-deep',
          'seaborn-v0_8-muted',
          'seaborn-v0_8-notebook',
          'seaborn-v0_8-paper',
          'seaborn-v0_8-pastel',
          'seaborn-v0_8-poster',
          'seaborn-v0_8-talk',
          'seaborn-v0_8-ticks',
          'seaborn-v0_8-white',
          'seaborn-v0_8-whitegrid',
          'tableau-colorblind10']
```

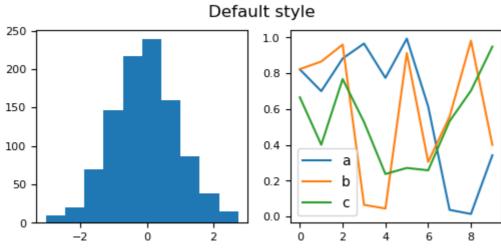
In [8]: for style in plt.style.available[:7] :
 with plt.style.context(style):
 hist_and_lines(f'{style=}')
hist_and_lines()







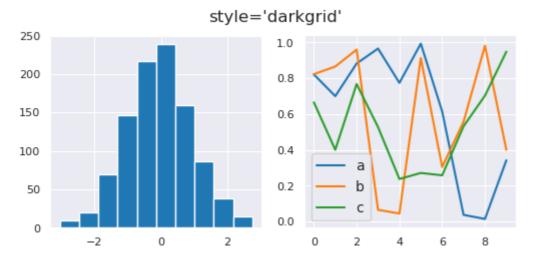


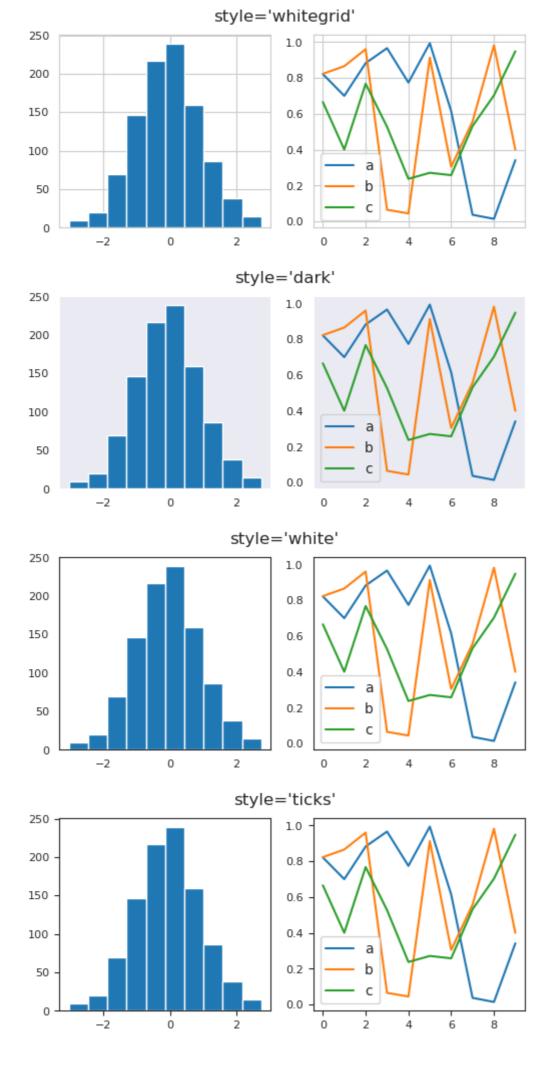


Seaborn Styles

- Five styles: darkgrid, whitegrid, dark, white, and ticks
- sns.set_style(stylename) → set global style
 - Use context manager to change the style locally:
 - with sns.axes_style(stylename):
 make_a_plot()

```
In [9]: for style in ['darkgrid', 'whitegrid', 'dark', 'white', 'ticks']:
    with sns.axes_style(style):
        hist_and_lines(f'{style=}')
    hist_and_lines()
```





Default style 1.0 200 150 100 50 -2 0.2 0.4 0.2 0.0 0.4 0.2 0.0 0.4 0.2 0.8

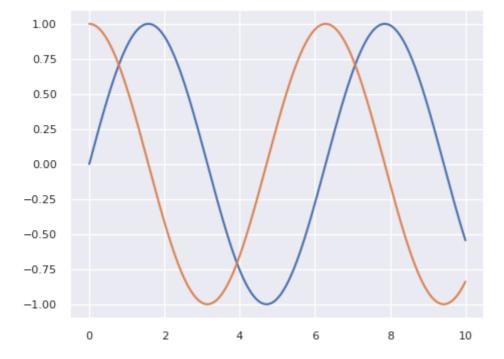
• sns.set() == sns.set_theme() → sns.set_style('darkgrid') and more

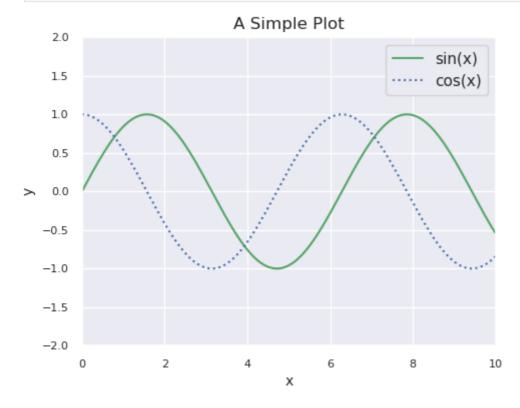
```
In [10]: sns.set()
    mpl.rcParams['figure.figsize'] = (5.33,4)
    mpl.rcParams['axes.labelsize'] = 10  # Example: 14 points
    mpl.rcParams['xtick.labelsize'] = 8  # Example: 12 points for x-axis ticks
    mpl.rcParams['ytick.labelsize'] = 8  # Example: 12 points for y-axis ticks
    hist_and_lines("New default style")
```



Line, Axes and Label properties

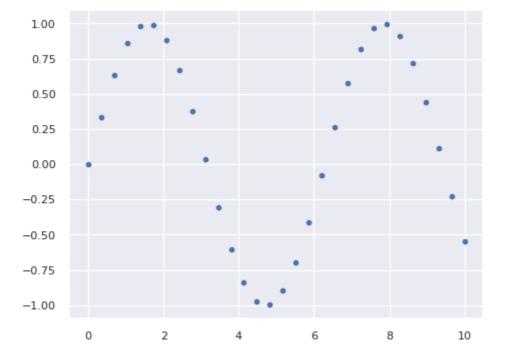
```
In [11]: x = np.linspace(0, 10, 1000)
    ax = plt.axes()
    ax.plot(x, np.sin(x))
    ax.plot(x, np.cos(x));
```



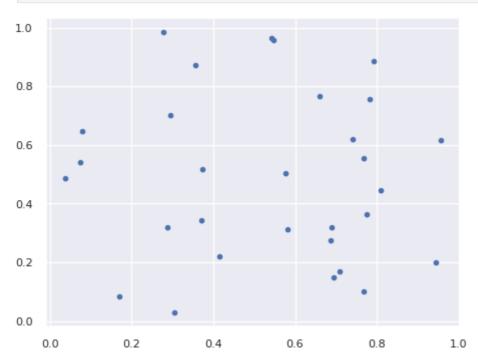


Scatter Plots

```
In [13]: x = np.linspace(0, 10, 30)
y = np.sin(x)
#plt.plot(x, y);  # default: '-' use lines
plt.plot(x, y, '.'); # use point as marker
```



```
In [14]: x,y = np.random.rand(2,30)
#plt.plot(x, y); # default: '-' use lines
plt.plot(x, y, '.'); # use point as marker
```



• scatter(x, y, s=, c=, cmap=) \rightarrow scatter plot with individual size and color

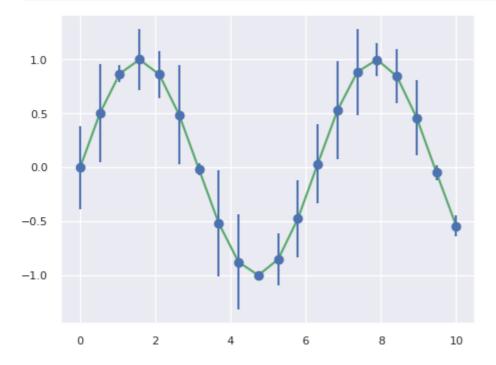
```
In [15]: x,y,sizes,colors = np.random.rand(4,30)
    plt.scatter(x, y, s=sizes*1000, alpha=0.3, c=colors, cmap='viridis')
    plt.colorbar(); # show color scale
```



Errorbars

• plt.errorbar(x, y, yerr=) \rightarrow an error bar plot

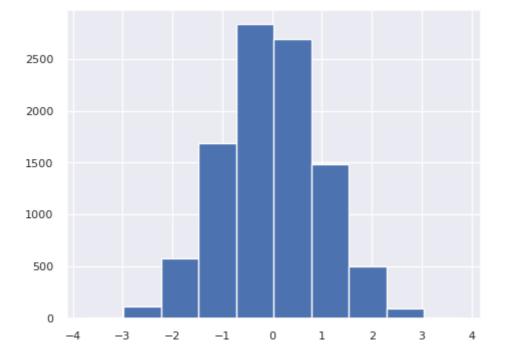
```
In [16]: x = np.linspace(0, 10, 20)
y = np.sin(x)
dy = 0.5 * np.random.rand(x.size)
plt.plot(x, np.sin(x), 'g')
plt.errorbar(x, np.sin(x), yerr=dy, fmt='o');
```



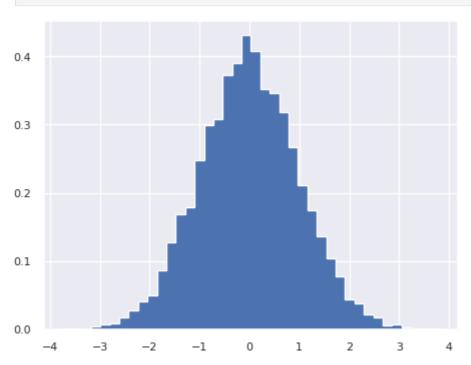
Histograms

• plt.hist(x, bins=, density=, histtype=) → a histogram plot

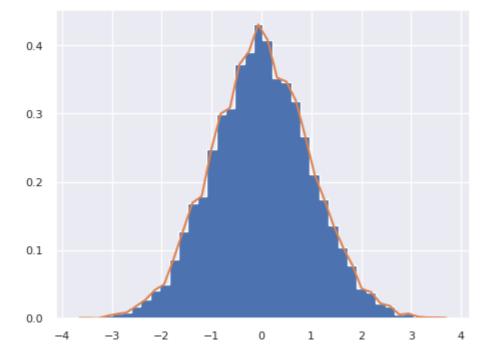
```
In [17]: x = np.random.randn(10000)
   plt.hist(x);
```



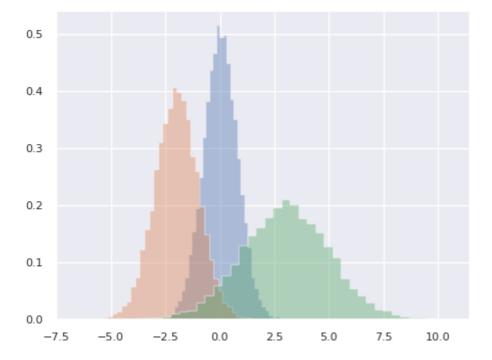
In [18]: plt.hist(x, bins=40, density=True, histtype='stepfilled');



In [19]: density, bins, patches = plt.hist(x, bins=40, density=True, histtype='stepfilled')
Calculate the center of the bins
bin_centers = (bins[:-1] + bins[1:]) / 2
Plot the probability density as a line
plt.plot(bin_centers, density);



```
In [20]: x1 = np.random.normal(0, 0.8, 10000)
    x2 = np.random.normal(-2, 1, 10000)
    x3 = np.random.normal(3, 2, 10000)
    kwargs = dict(bins=40, density=True, histtype='stepfilled', alpha=0.4)
    plt.hist(x1, **kwargs)
    plt.hist(x2, **kwargs);
    plt.hist(x3, **kwargs);
```



Seaborn

- Matplotlib predated Pandas by more than a decade
 - It is not designed for use with Pandas (is being updated)
- Seaborn provides an API on top of Matplotlib and integrates with Pandas
 - Statistical tools: regressions, errors and distributions
 - Simple multivariate visualization

```
In [21]: a = np.random.multivariate_normal(mean=[0, 1], cov=[[5, 2], [2, 2]], size=2000)
df = pd.DataFrame(a, columns=['x', 'y'])
df.head()
```

```
        x
        y

        0
        0.422732
        0.152363

        1
        -0.733416
        0.678723

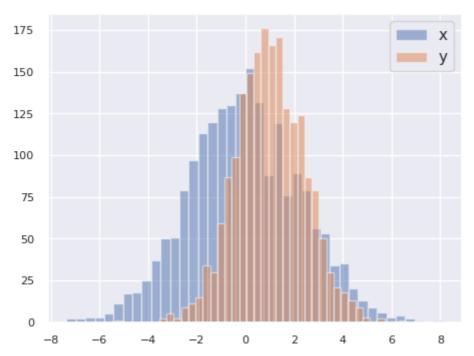
        2
        -1.825696
        0.841521

        3
        2.113652
        1.340771

        4
        1.254496
        1.035458
```

Plotting two histograms (widths of the bins differ)

```
In [22]: plt.hist(df['x'], bins=40, alpha=0.5, label='x')
    plt.hist(df['y'], bins=40, alpha=0.5, label='y')
    plt.legend();
```



Seaborn can plot the histograms of the DataFrame variables (equal bin widths):

In [23]: sns.histplot(df, bins=40);



Kernel density estimation of the DataFrame variables:

Two-dimensional visualization of the kernel density estimation:

-2.5

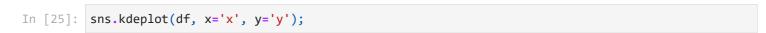
0.0

0.02

0.00

-7.5

-5.0

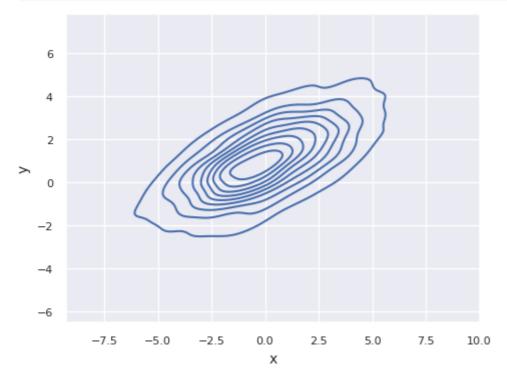


2.5

5.0

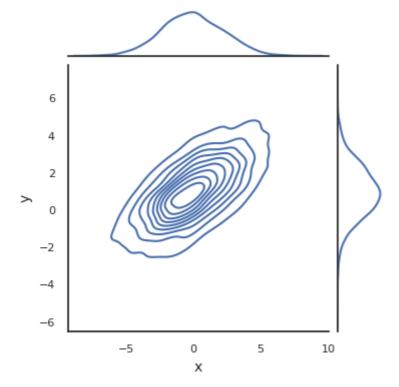
10.0

7.5

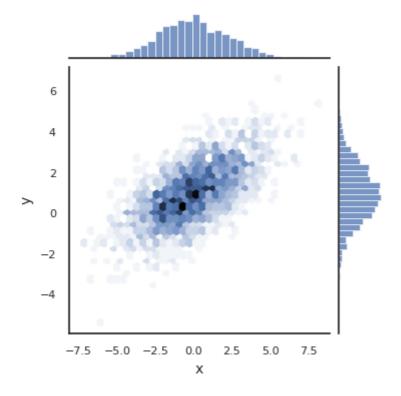


Joint and marginal distributions together:

```
In [26]: with sns.axes_style('white'):
    sns.jointplot(df, x='x', y='y', kind='kde', height=4);
```



Same but using an hexagonally based histogram instead:



Multivariate visualization

Iris dataset: measurements of petals and sepals of three iris species:

| | sepal_length | sepal_width | petal_length | petal_width | species |
|---|--------------|-------------|--------------|-------------|---------|
| 0 | 5.1 | 3.5 | 1.4 | 0.2 | setosa |
| 1 | 4.9 | 3.0 | 1.4 | 0.2 | setosa |
| 2 | 4.7 | 3.2 | 1.3 | 0.2 | setosa |
| 3 | 4.6 | 3.1 | 1.5 | 0.2 | setosa |
| 4 | 5.0 | 3.6 | 1.4 | 0.2 | setosa |

The correlation matrix of the data:

```
In [29]: iris.drop(['species'], axis=1).corr()
```

| | sepal_length | sepal_width | petal_length | petal_width |
|--------------|--------------|-------------|--------------|-------------|
| sepal_length | 1.000000 | -0.117570 | 0.871754 | 0.817941 |
| sepal_width | -0.117570 | 1.000000 | -0.428440 | -0.366126 |
| petal_length | 0.871754 | -0.428440 | 1.000000 | 0.962865 |
| petal_width | 0.817941 | -0.366126 | 0.962865 | 1.000000 |

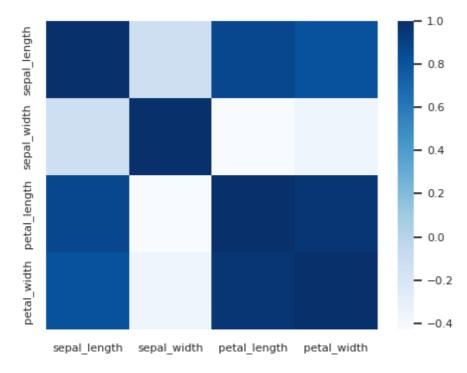
A heat map of the correlation matrix:

```
In [30]: c = iris.drop(['species'], axis=1).corr()
sns.heatmap(c, cmap="Blues")
```



Out[28]:

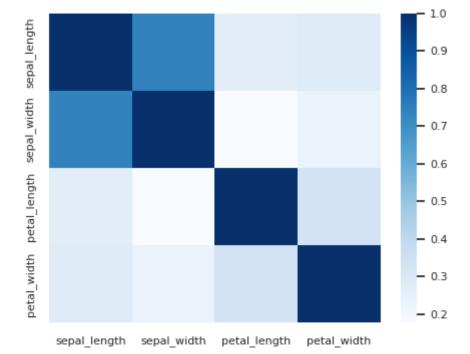
Out[29]:



A heat map of the correlation matrix just for setosa specie:

```
In [31]: c = iris[iris['species']=='setosa'].drop('species', axis=1).corr()
    sns.heatmap(c, cmap="Blues")
```

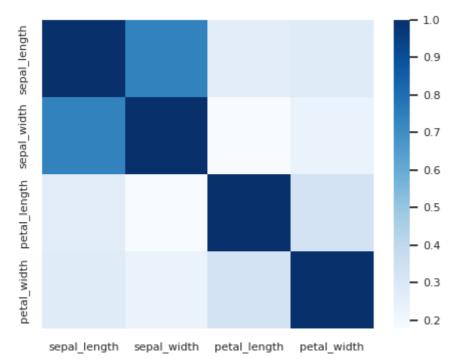
Out[31]: <Axes: >



Same but using the species variable as index:

```
In [32]: c = iris.set_index('species').loc['setosa'].corr()
sns.heatmap(c, cmap="Blues")
```

Out[32]: <Axes: >



Correlation matrix for each specie:

```
In [33]: iris.groupby('species').corr()
```

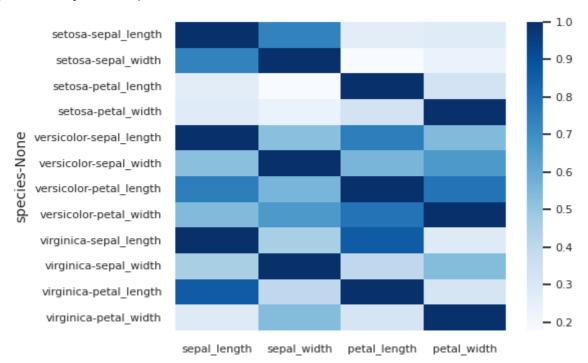
| Out[33]: sepal_len | ngth | sepal_width | petal_length | petal_width |
|--------------------|------|-------------|--------------|-------------|
|--------------------|------|-------------|--------------|-------------|

| species | | | | | |
|------------|--------------|----------|----------|----------|----------|
| setosa | sepal_length | 1.000000 | 0.742547 | 0.267176 | 0.278098 |
| | sepal_width | 0.742547 | 1.000000 | 0.177700 | 0.232752 |
| | petal_length | 0.267176 | 0.177700 | 1.000000 | 0.331630 |
| | petal_width | 0.278098 | 0.232752 | 0.331630 | 1.000000 |
| versicolor | sepal_length | 1.000000 | 0.525911 | 0.754049 | 0.546461 |
| | sepal_width | 0.525911 | 1.000000 | 0.560522 | 0.663999 |
| | petal_length | 0.754049 | 0.560522 | 1.000000 | 0.786668 |
| | petal_width | 0.546461 | 0.663999 | 0.786668 | 1.000000 |
| virginica | sepal_length | 1.000000 | 0.457228 | 0.864225 | 0.281108 |
| | sepal_width | 0.457228 | 1.000000 | 0.401045 | 0.537728 |
| | petal_length | 0.864225 | 0.401045 | 1.000000 | 0.322108 |
| | petal_width | 0.281108 | 0.537728 | 0.322108 | 1.000000 |

Correlation matrix heat map for each specie:



Out[34]: <Axes: ylabel='species-None'>



Plot with pair-wise scatter plots and kde:

```
In [35]: sns.pairplot(iris, hue='species', height=1.5);
```

