

# Introduction to Python: Seaborn

NYC Data Science Academy

# **OVERVIEW**

- Seaborn overview
- Univariate plots
- Bivariate plots
- Multivariate plots
- Advanced plots

### **Visualization in Python**

This course covers two parts: matplotlib and seaborn. This slide covers the package Seaborn.

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Seaborn is a Python visualization library based on matplotlib. It provides a high-level interface for drawing attractive statistical graphics. Here is the <u>documentation</u>.

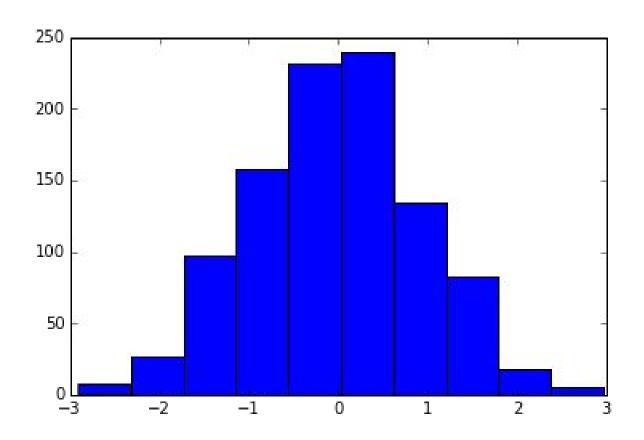
We first import the packages matplotlib.pyplot and visualize with it.

```
%pylab inline import matplotlib.pyplot as plt
```

The package Pandas is imported for data manipulation, Numpy and SciPy is imported to generate random numbers and distribution.

```
from scipy import stats
import pandas as pd
import numpy as np
```

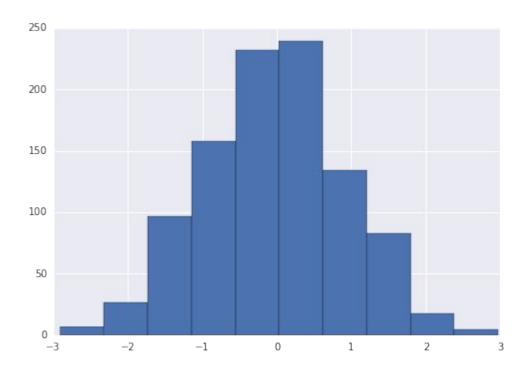
The code below visualize a normal distribution.





Import seaborn and visualize again:

import seaborn as sns
plt.hist(df.norm)





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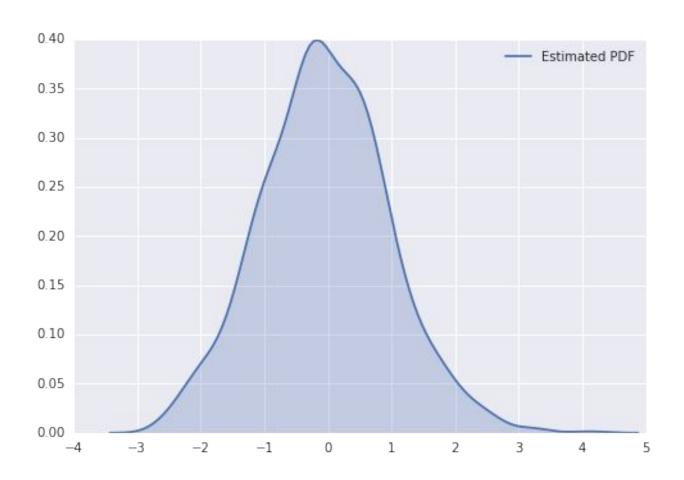
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We generate some arrays that we will visualize.

```
np.random.seed(2)
x = np.random.randn(1e3)
y = 2*x + np.random.rand(1e3)
```

We can of course visualize the distribution of x with histogram. However, seaborn provides a nice function that smooths out the histogram to estimate the distribution.

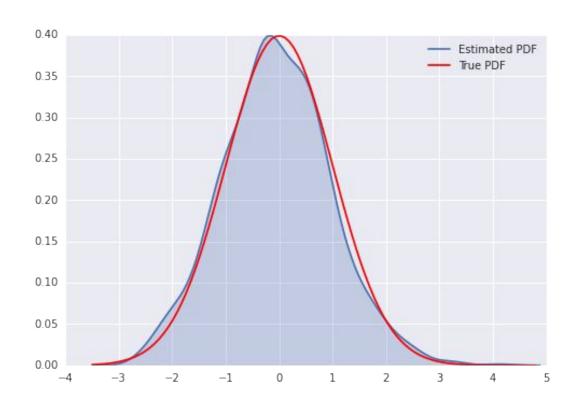
```
sns.kdeplot(x, shade=True, label='Estimated PDF')
```





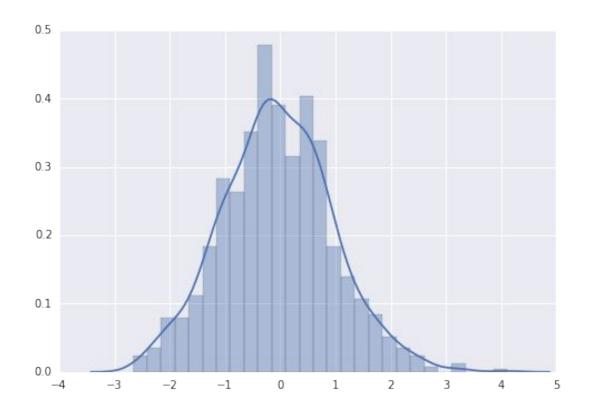
### **Exercise**

Modify the code above to obtain the plot below:





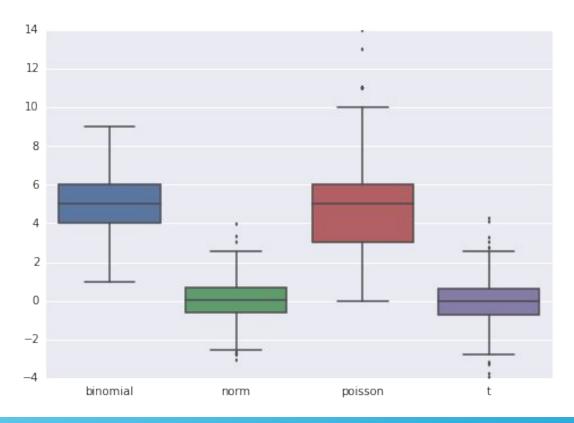
It is possible to combine histogram and the distribution estimate plot:



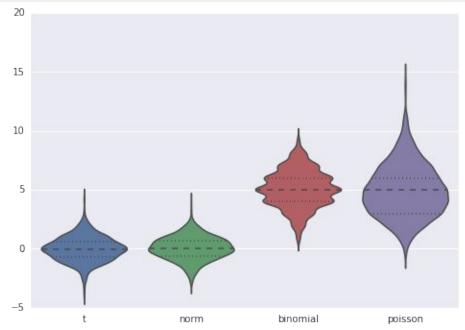


We may also visualize the distribution of multiple features in one plot:

sns.boxplot(df)



• We see that seaborn understand data frames. Here we also demonstrate how to order columns in a data frame.

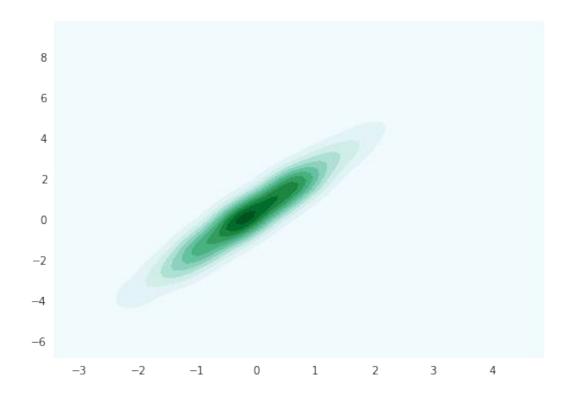


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### **Bivariate plot**

The kdeplot() function can be used to plot 2D density.





### **Exercise**

Sketch the 2D density plot for the normal random sample and the binomial random sample in our df.

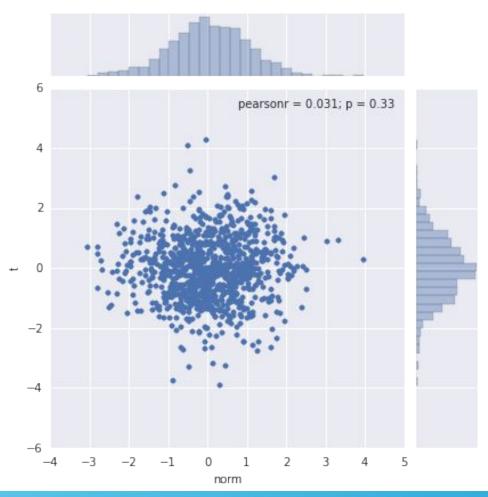
### **Bivariate plot**

The joinplot() function combines histogram and scatter plot.

```
sns.jointplot(df.norm, df.t)
```

❖ Below we see "pearsonr=0.031" which indicates the pearson correlation of these two variables. We also see "p=0.33", which indicates that there is no significant linear relation between the two variables.

# **Bivariate plot**





#### **Exercise**

- ❖ Visualize the variables x and y we created in the previous slide. Explain the pearsonr and p for this case.
- Visualize with the same function as above. This time specify kind="hex", kind="reg" and kind="kde", respectively.

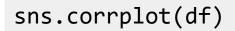


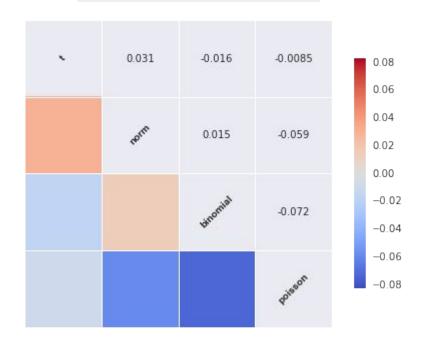
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### Multivariate plot

A scatter plot or a joint plot just show the correlation of two variables, corrplot is similar to scatter-matrix, it shows the correlations between multiple variables instead of scatters.







### Multivariate plot

Seaborn also provides sample datasets.

```
titanic = sns.load_dataset("titanic").dropna()
```

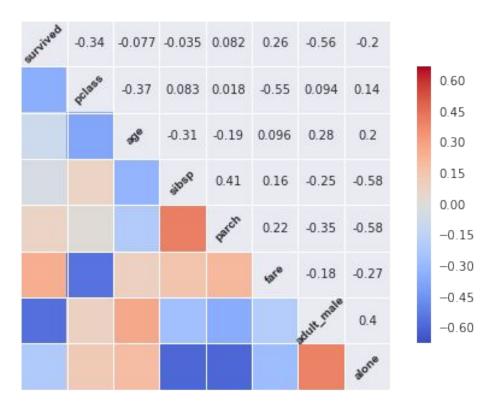
This is a famous dataset which was originally used to demonstrate a classification problem: using the record of each passengers in the ship to predict the probability that he or she survived. Run the code below to see the data frame.

```
titanic.head()
```

### **Multivariate plot**

#### Plot the correlation:

sns.corrplot(titanic)





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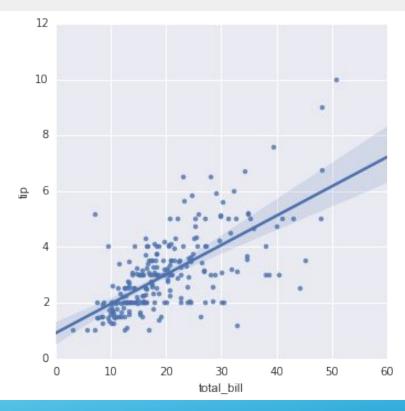
### **Advanced plot**

Seaborn also provides another famous sample dataset:

```
tips = sns.load_dataset("tips")
tips.head()
```

This is a famous dataset record the information of people of different sex, being a smoker or not, visiting at different day or for different meal, the size of parties they joined, and the tip and the total bill they pay.

To visualize the relation between total\_bill and tip, of course we can use scatter plot. However, seaborn allows fitting with linear model as well.

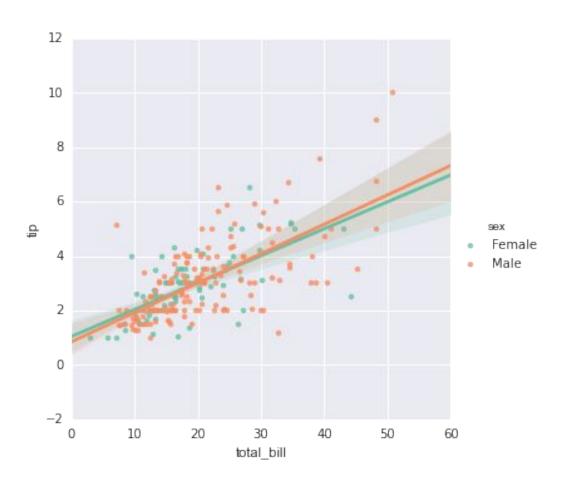




We can further split the data into "Male" and "Female" parts and visualize them.

```
sns.lmplot("total_bill", "tip", tips,\
hue="sex", palette="Set2");
```

- Here we used:
  - hue indicates according to which column we group our data.
  - palette simply specifies the color we want to use.

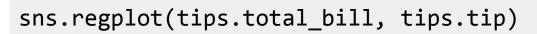


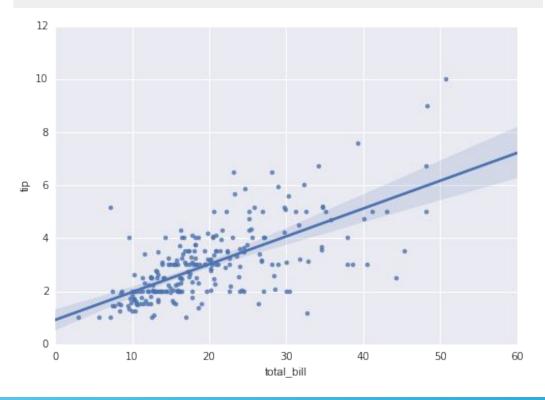


### **Exercise**

\* Make the same visualization as in the previous slide. This time group the data according to that if the observation is a smoker or not.

Implot only supports data frames, regplot also accepts data passed directly as numpy arrays or pandas series objects.

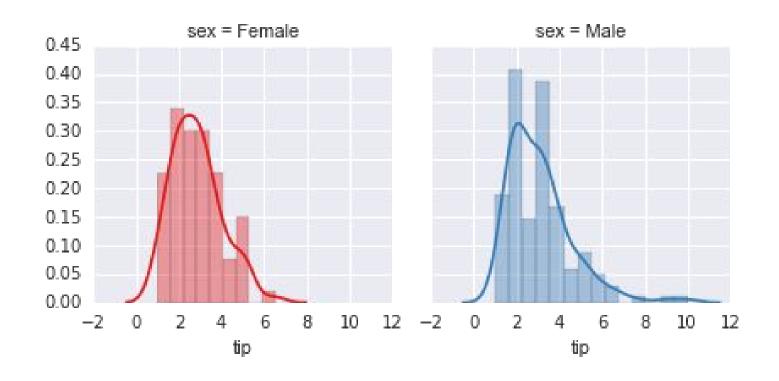




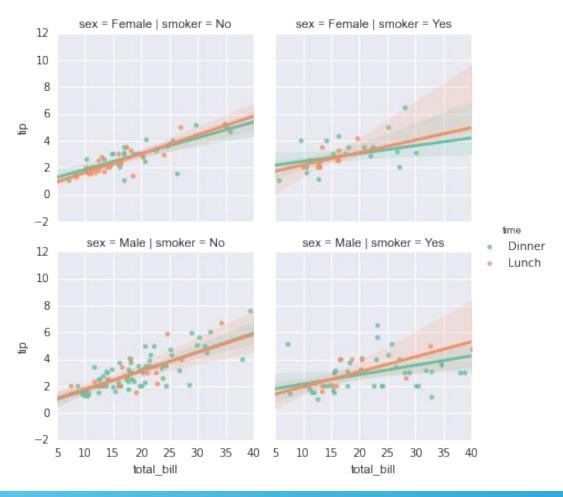


• We often need to compare the same kind of plot for different features.
Functions for faceting comes in handy.

Within the function FacetGrid(), hue indicates sketching a separate plot for each sex; col specifies that each plot is placed in a new column; palette specifies the colors. Then we use .map() method to specify the type of the plot and the feature we want to visualize.



Facet can be used to differentiate multiple factors as well.



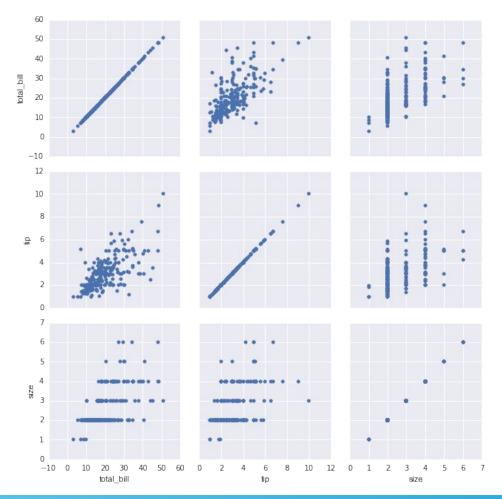


### Advanced plot: pairwise plot

The function FacetGrid help you explore the specific variables conditioned on different levels. Function PairGrid is useful to explore the relationships between pairs of variables.

```
tipGrid = sns.PairGrid(tips)
tipGrid.map(plt.scatter)
```

# Advanced plot: pairwise plot





#### **Exercise 5**

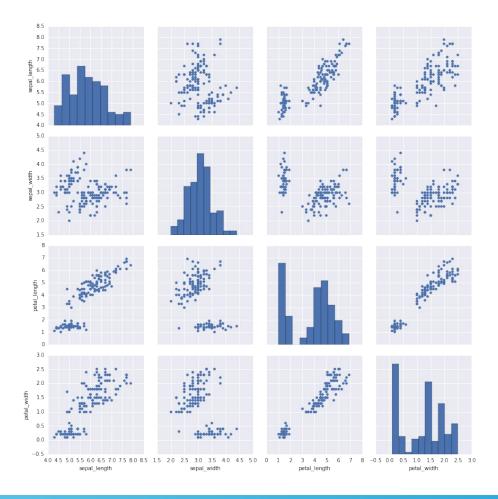
Download the iris data frame built in the seaborn package.

```
iris = sns.load_dataset("iris")
iris.head()
```

Then pair the four numeric features and sketch scatter plot for each pair.

Some examples with different type of plot and colors.

```
g = sns.PairGrid(iris)
g.map_diag(plt.hist)
g.map_offdiag(plt.scatter)
```





Some examples with different type of plot and colors.

