

```
In [1]: import seaborn as sns
import matplotlib.pyplot as plt

plt.style.use('ggplot')
```

Categorical Plots



Categorical Scatter Plot

- Stripplot
- Swarmplot

Categorical Distribution Plots

- Boxplot
- Violinplot

Categorical Estimate Plot -> for central tendency

- Barplot
- Pointplot
- Countplot

Figure level function -> catplot

```
In [2]: # import datasets

tips = sns.load_dataset('tips')
iris = sns.load_dataset('iris')
```

```
In [3]: tips.sample(2)
```

Out[3]:

| | total_bill | tip | sex | smoker | day | time | size |
|----|------------|------|--------|--------|------|--------|------|
| 94 | 22.75 | 3.25 | Female | No | Fri | Dinner | 2 |
| 88 | 24.71 | 5.85 | Male | No | Thur | Lunch | 2 |

```
In [4]: iris.sample(2)
```

```
Out[4]:
```

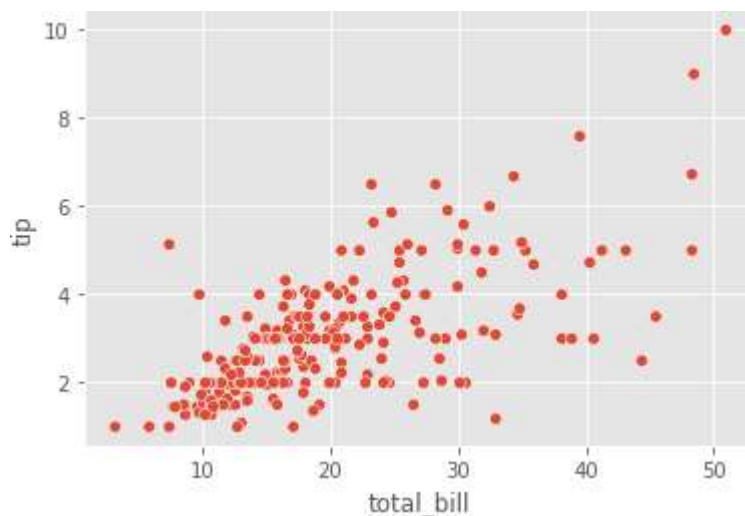
| | sepal_length | sepal_width | petal_length | petal_width | species |
|-----|--------------|-------------|--------------|-------------|------------|
| 53 | 5.5 | 2.3 | 4.0 | 1.3 | versicolor |
| 103 | 6.3 | 2.9 | 5.6 | 1.8 | virginica |

Categorical Scatter Plots

```
In [5]: # ON Numerical data
```

```
sns.scatterplot(data=tips, x='total_bill', y='tip')
```

```
Out[5]: <AxesSubplot:xlabel='total_bill', ylabel='tip'>
```



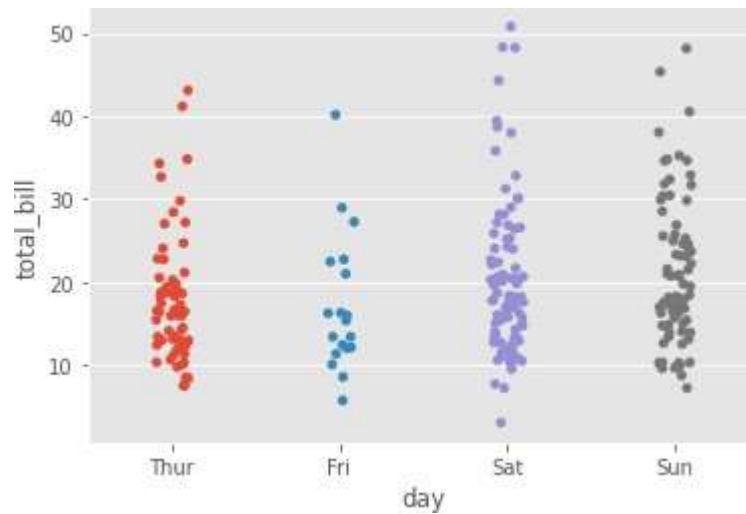
1.strip plot

In [6]: *# axes level function*

On categorical data

```
sns.stripplot(data=tips,x='day',y='total_bill')
```

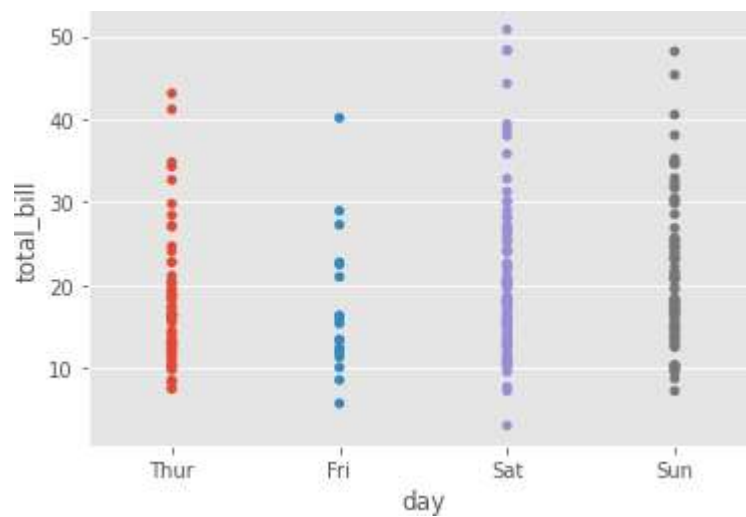
Out[6]: <AxesSubplot:xlabel='day', ylabel='total_bill'>



In [7]: *# jitter*

```
sns.stripplot(data=tips,x='day',y='total_bill', jitter=False)
```

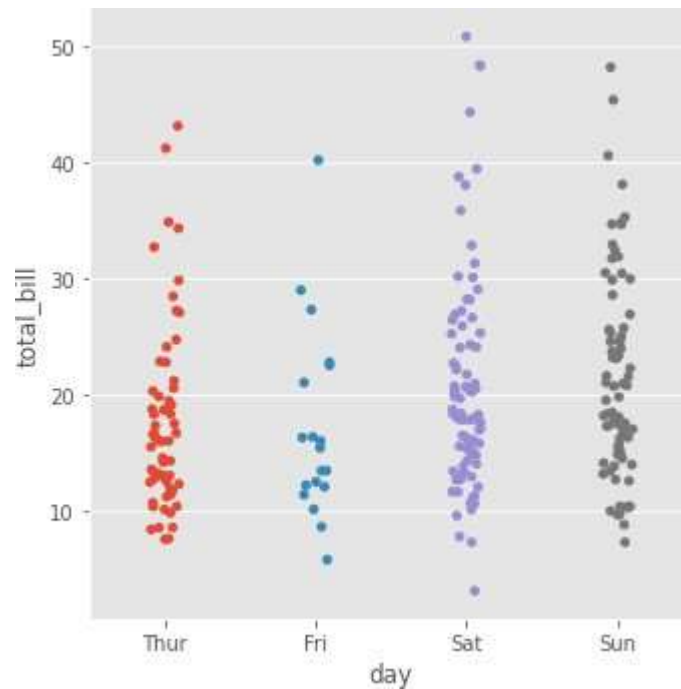
Out[7]: <AxesSubplot:xlabel='day', ylabel='total_bill'>



catplot

```
In [8]: # catplot -> Categorical plot  
  
# Figure Level functions  
  
sns.catplot(data=tips,x='day',y='total_bill',kind = 'strip')
```

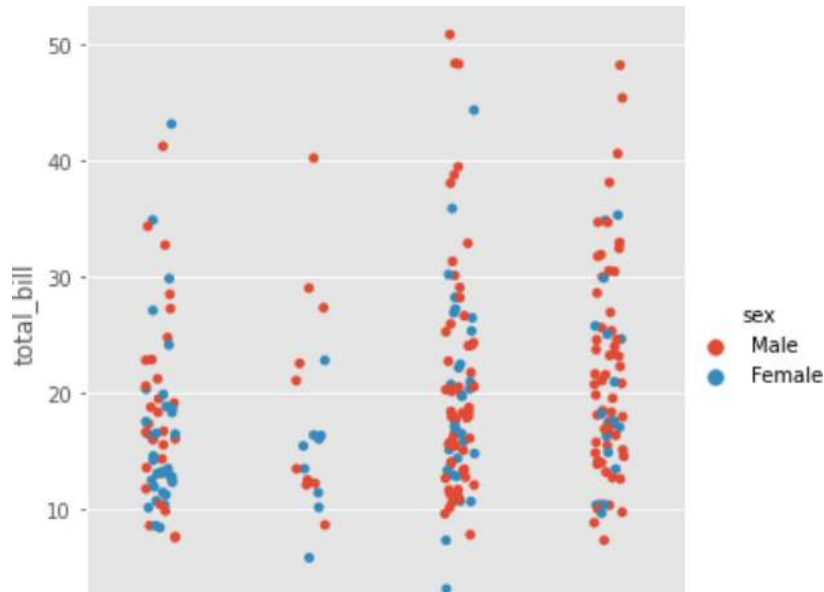
Out[8]: <seaborn.axisgrid.FacetGrid at 0x1e85ea0ed90>



In [9]: `# hue`

```
sns.catplot(data=tips,x='day',y='total_bill',kind = 'strip',  
            hue = 'sex')
```

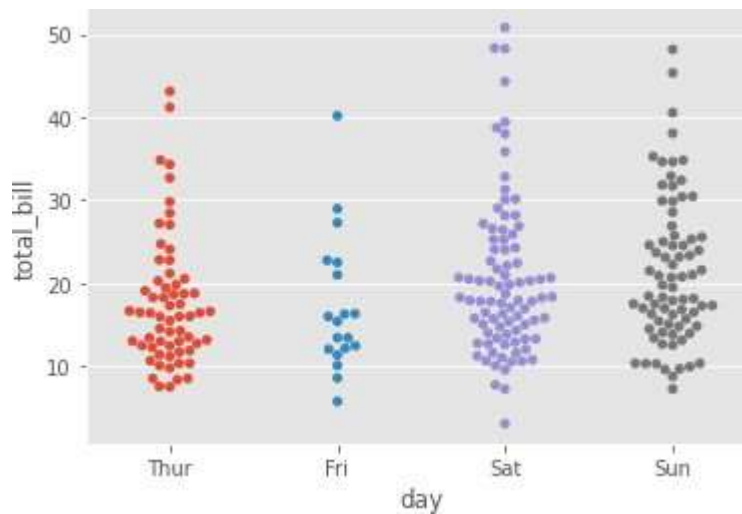
Out[9]: `<seaborn.axisgrid.FacetGrid at 0x1e85f320e50>`



2 .Swarm plot

In [10]: `sns.swarmplot(data=tips,x = 'day', y='total_bill')`

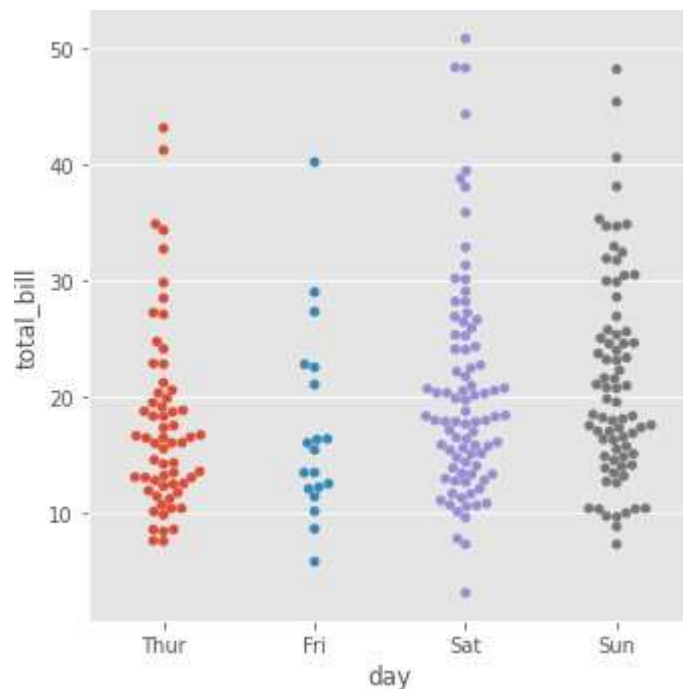
Out[10]: `<AxesSubplot:xlabel='day', ylabel='total_bill'>`



```
In [11]: # use by catplot
```

```
sns.catplot(data=tips,x ='day', y='total_bill',kind ='swarm')
```

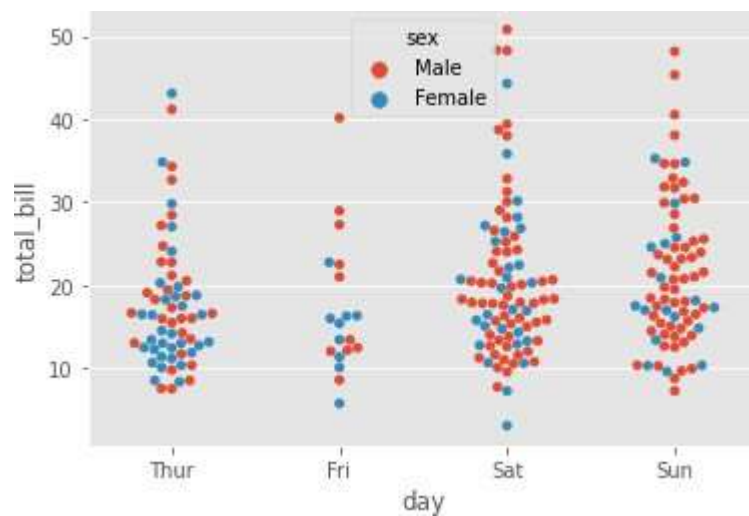
```
Out[11]: <seaborn.axisgrid.FacetGrid at 0x1e85f433850>
```



```
In [12]: # hue
```

```
sns.swarmplot(data=tips,x ='day', y='total_bill',hue='sex')
```

```
Out[12]: <AxesSubplot:xlabel='day', ylabel='total_bill'>
```

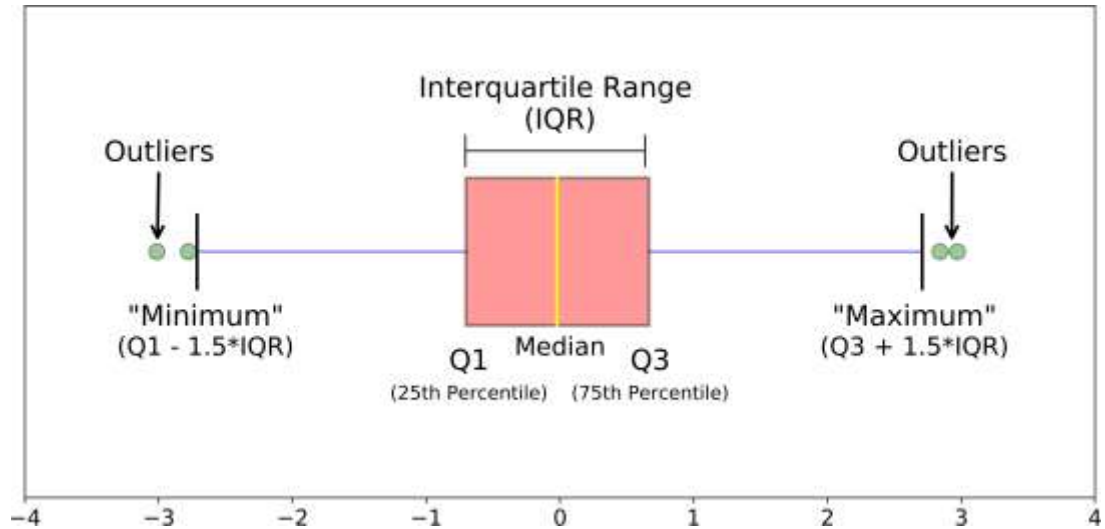


Categorical Distribution Plots

1. Boxplot

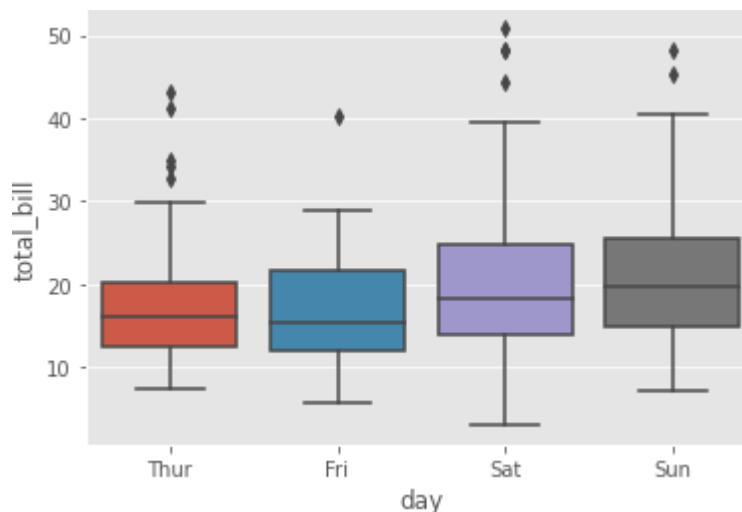
A boxplot is a standardized way of displaying the distribution of data based on a five number summary (“minimum”, first quartile [Q1], median, third quartile [Q3] and “maximum”).

- It can tell you about your outliers and what their values are.
- Boxplots can also tell you if your data is symmetrical,
- how tightly your data is grouped and if and how your data is skewed.



```
In [13]: # Box plot
sns.boxplot(data=tips,x='day',y='total_bill')
```

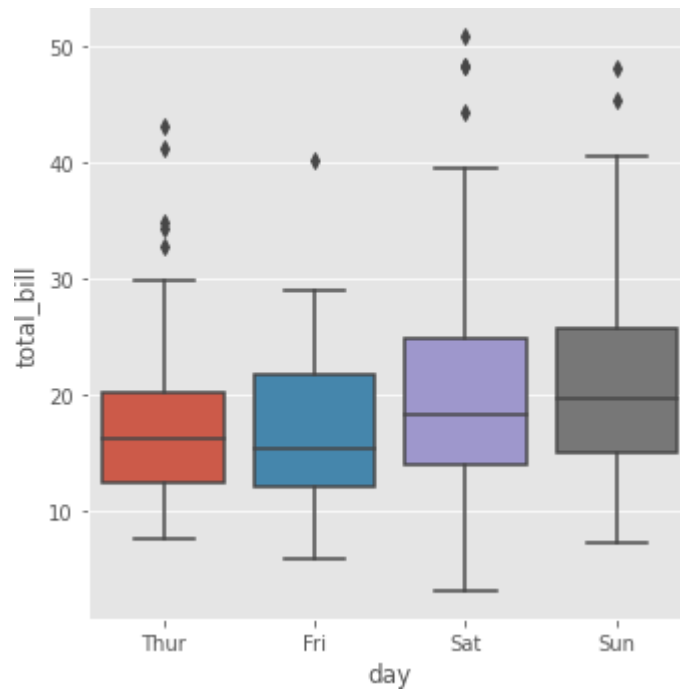
```
Out[13]: <AxesSubplot:xlabel='day', ylabel='total_bill'>
```



```
In [14]: # Using catplot - Figure Level
```

```
sns.catplot(data=tips,x='day',y='total_bill',kind='box')
```

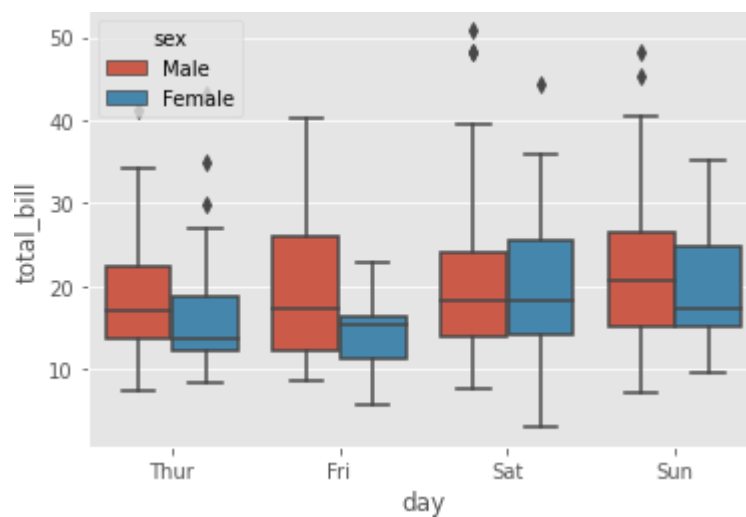
```
Out[14]: <seaborn.axisgrid.FacetGrid at 0x1e85f3b5dc0>
```



```
In [15]: # hue
```

```
sns.boxplot(data=tips,x='day',y='total_bill',hue='sex')
```

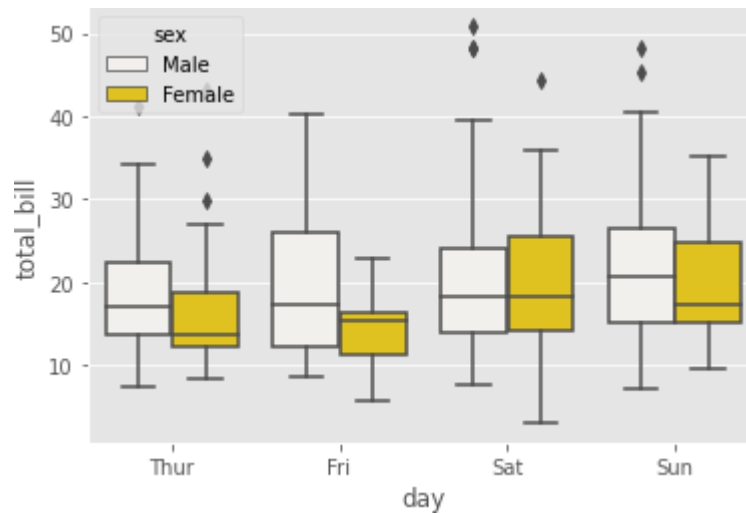
```
Out[15]: <AxesSubplot:xlabel='day', ylabel='total_bill'>
```




```
In [16]: # color
```

```
sns.boxplot(data=tips,x='day',y='total_bill',hue='sex', color = 'gold')
```

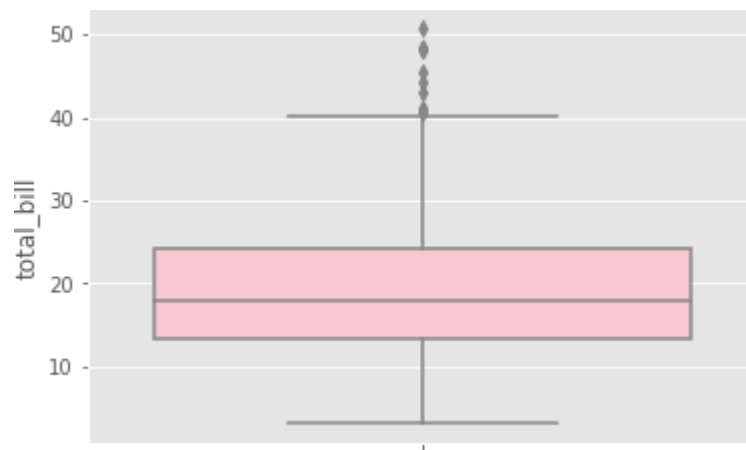
```
Out[16]: <AxesSubplot:xlabel='day', ylabel='total_bill'>
```



```
In [17]: # single boxplot -> numerical col
```

```
sns.boxplot(data=tips , y='total_bill', color ='pink')
```

```
Out[17]: <AxesSubplot:ylabel='total_bill'>
```

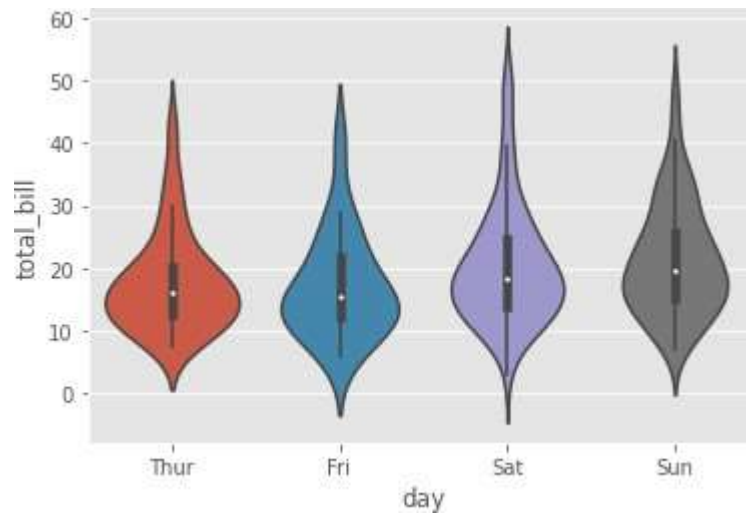


2 . Voilin Plot

```
In [18]: # Violinplot = (Boxplot + KDEplot)

sns.violinplot(data=tips,x='day',y='total_bill')
```

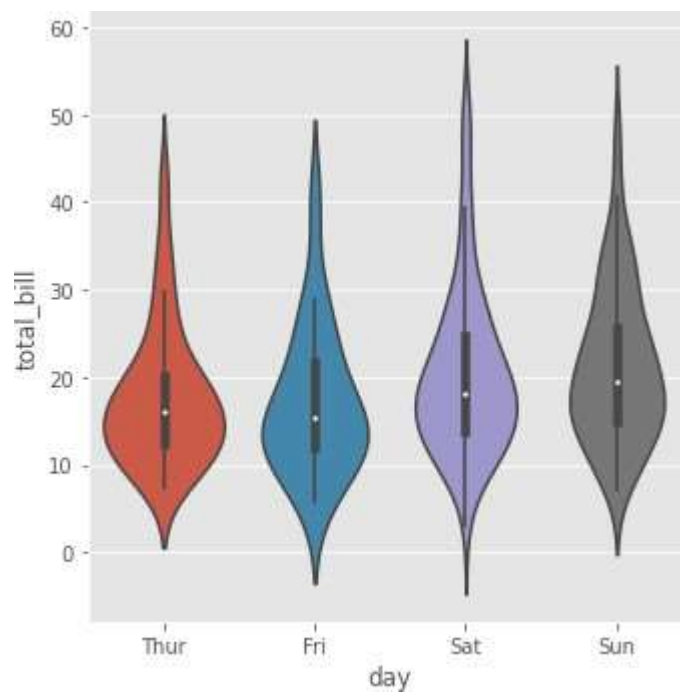
Out[18]: <AxesSubplot:xlabel='day', ylabel='total_bill'>



```
In [19]: # catplot -figure level function

sns.catplot(data=tips,x='day',y='total_bill',kind='violin')
```

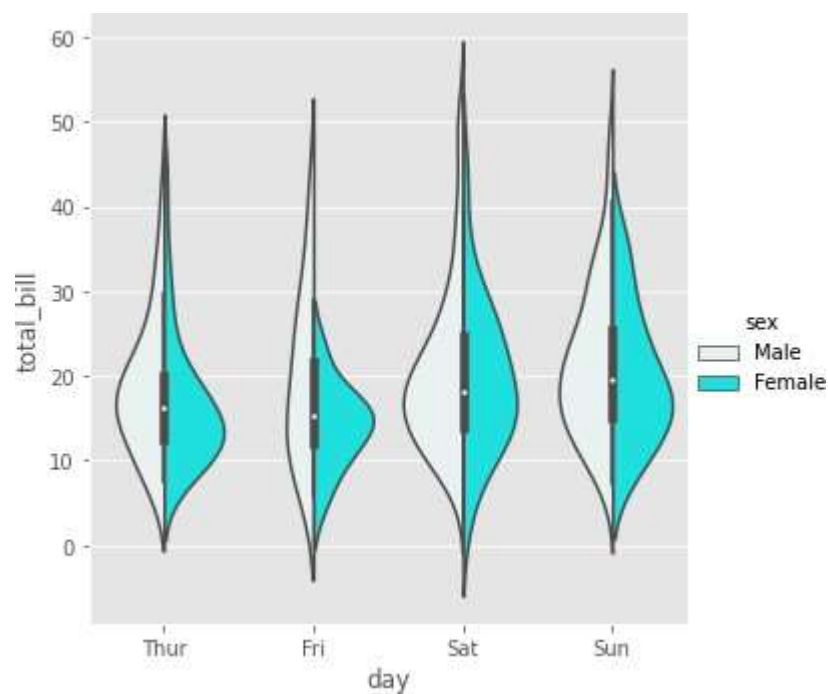
Out[19]: <seaborn.axisgrid.FacetGrid at 0x1e85f61a490>



In [20]: `# Color`

```
sns.catplot(data=tips,x='day',y='total_bill',  
            kind='violin', hue='sex', split=True, color='cyan')
```

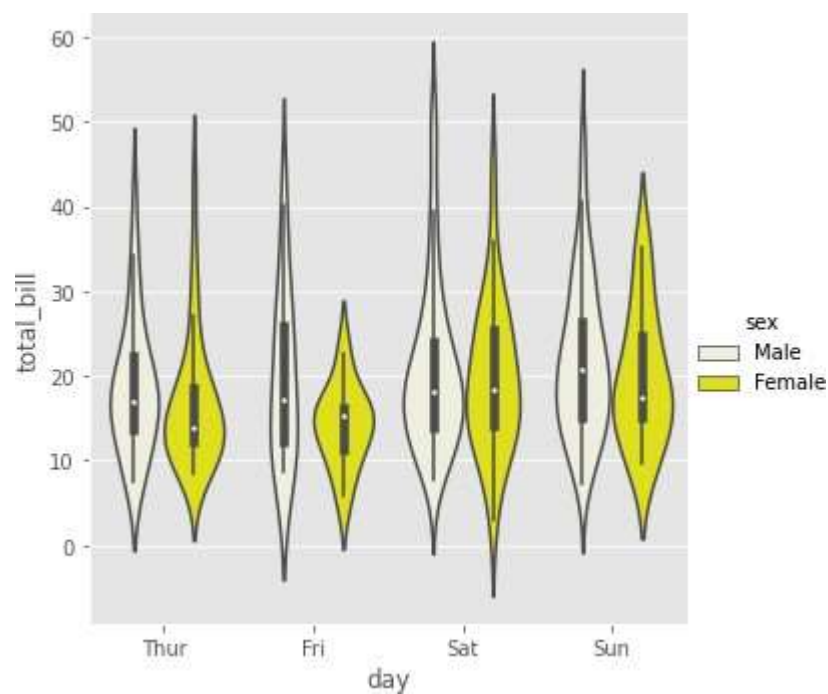
Out[20]: `<seaborn.axisgrid.FacetGrid at 0x1e85f5b2580>`



In [21]: # hue

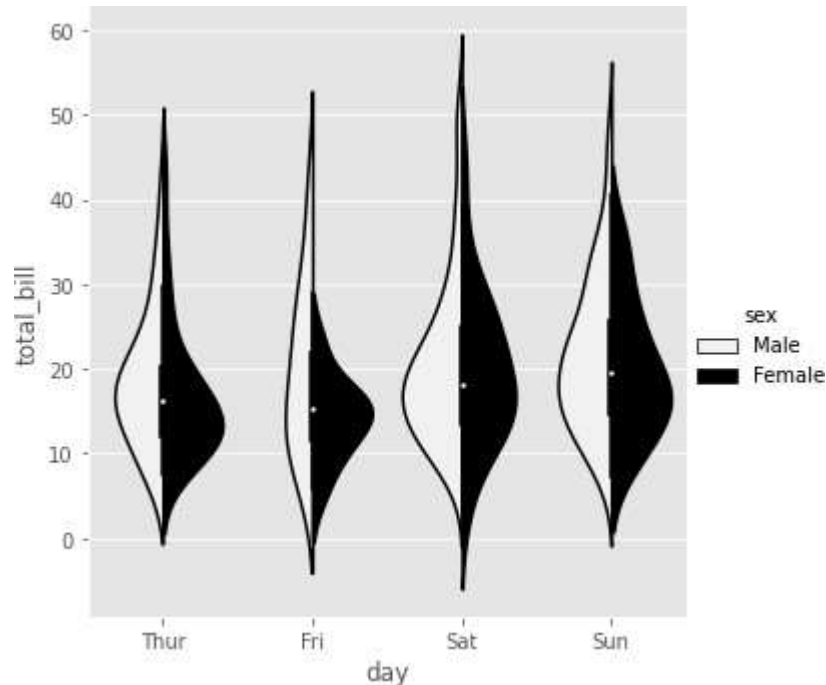
```
sns.catplot(data=tips,x='day',y='total_bill',kind='violin',  
            hue='sex',color='yellow')
```

Out[21]: <seaborn.axisgrid.FacetGrid at 0x1e860a5da60>



```
In [22]: # split
sns.catplot(data=tips,x='day',y='total_bill',
            kind='violin', hue='sex',
            color='black', split=True)
```

Out[22]: <seaborn.axisgrid.FacetGrid at 0x1e860a0f580>



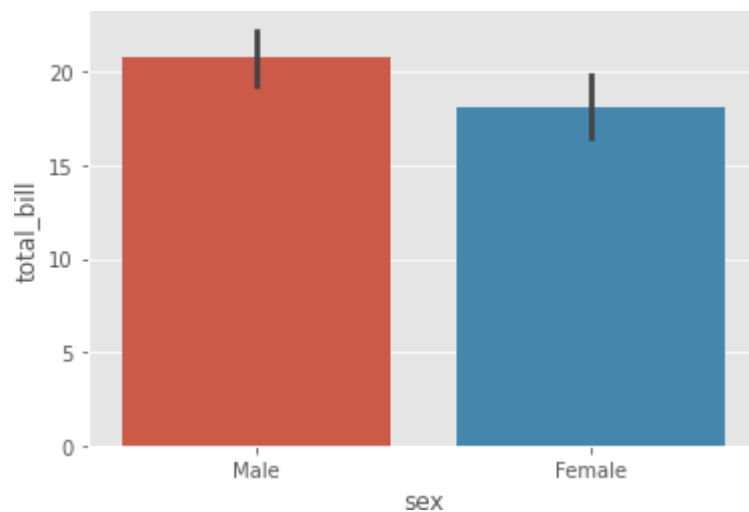
Categorical Estimate Plot

1.barplot

- When there are multiple observations in each category, it also uses bootstrapping to compute a confidence interval around the estimate, which is plotted using **error bars**

```
In [23]: sns.barplot(data=tips, x='sex', y='total_bill')
```

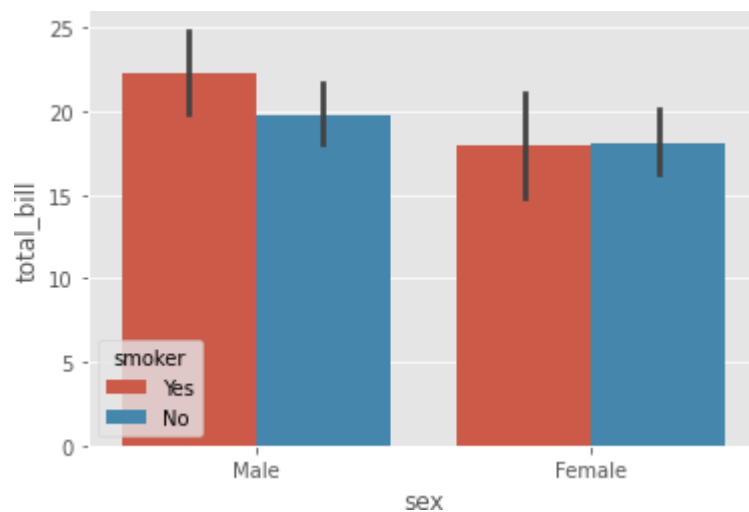
```
Out[23]: <AxesSubplot:xlabel='sex', ylabel='total_bill'>
```



```
In [24]: # hue
```

```
sns.barplot(data=tips, x='sex', y='total_bill', hue='smoker')
```

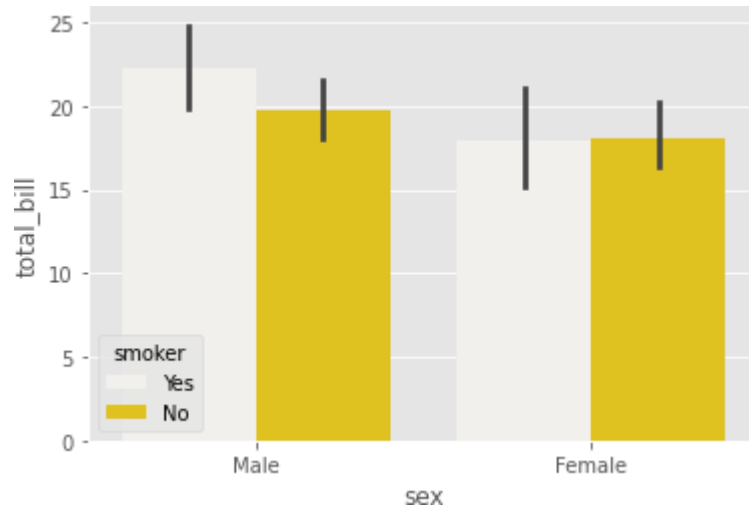
```
Out[24]: <AxesSubplot:xlabel='sex', ylabel='total_bill'>
```



In [25]: `# color`

```
sns.barplot(data=tips, x='sex', y='total_bill', hue='smoker',  
            color='gold')
```

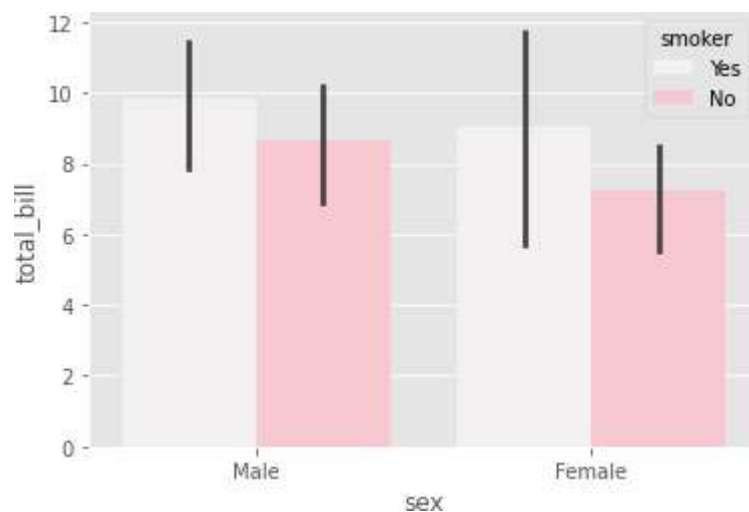
Out[25]: `<AxesSubplot:xlabel='sex', ylabel='total_bill'>`



In [26]: `# estimator`

```
import numpy as np  
  
sns.barplot(data=tips, x='sex', y='total_bill', hue='smoker', color='pink',  
            estimator=np.std)
```

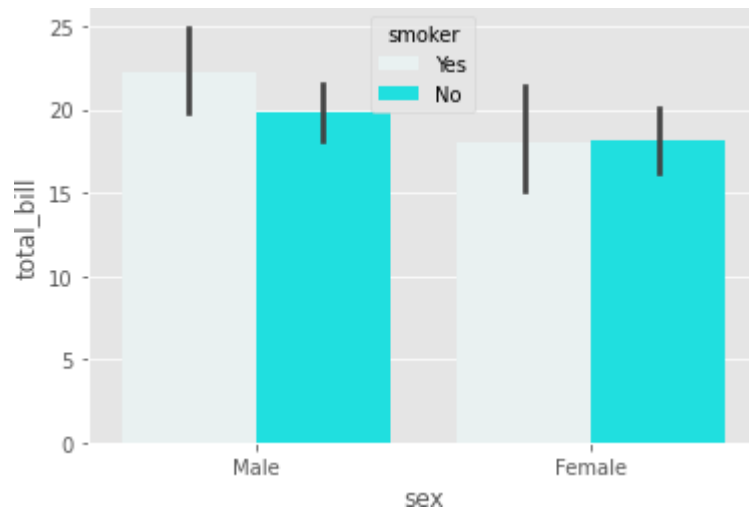
Out[26]: `<AxesSubplot:xlabel='sex', ylabel='total_bill'>`



```
In [27]: # estimator ( mean)
```

```
sns.barplot(data=tips, x='sex', y='total_bill',hue='smoker',color = 'cyan',  
            estimator=np.mean)
```

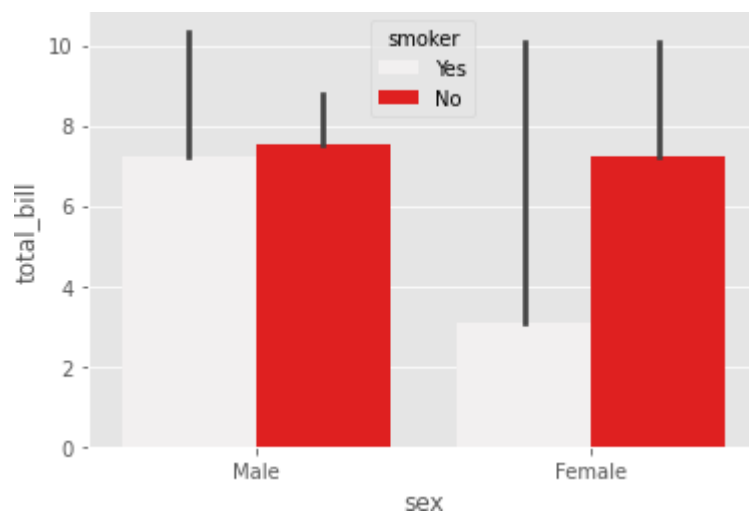
```
Out[27]: <AxesSubplot:xlabel='sex', ylabel='total_bill'>
```



```
In [28]: # estimator ( min)
```

```
sns.barplot(data=tips, x='sex', y='total_bill',hue='smoker',color = 'red',  
            estimator=np.min)
```

```
Out[28]: <AxesSubplot:xlabel='sex', ylabel='total_bill'>
```

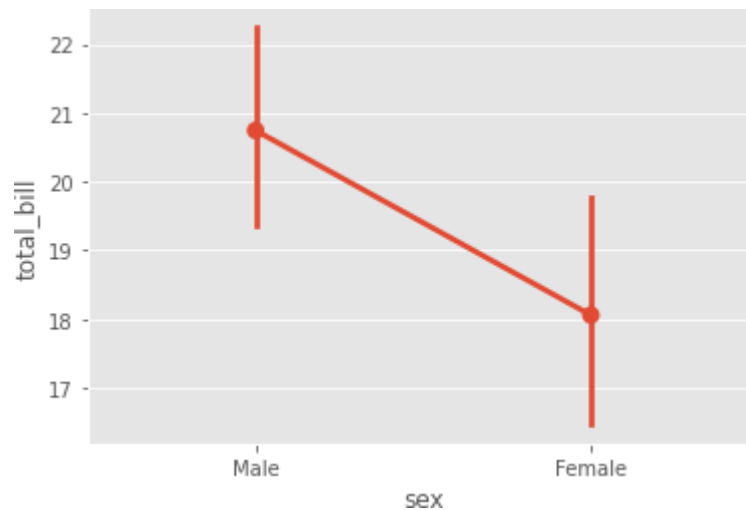


2. Point plot

A point plot represents an estimate of central tendency for a numeric variable by the position of

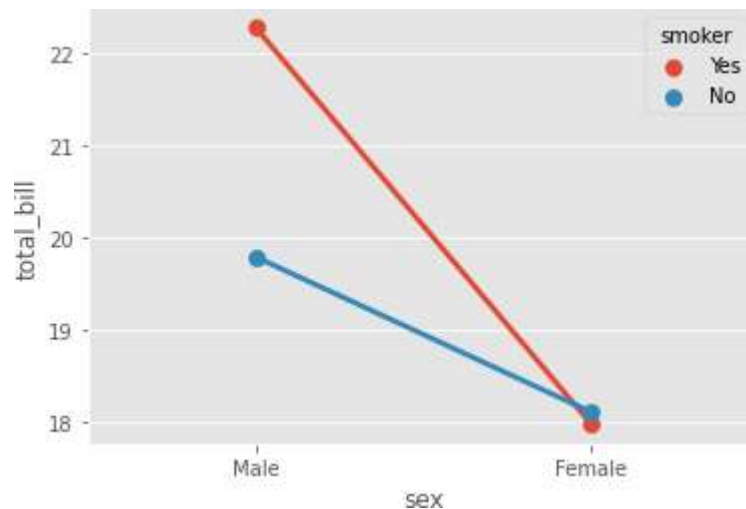

```
In [29]: sns.pointplot(data=tips, x='sex', y='total_bill')
```

```
Out[29]: <AxesSubplot:xlabel='sex', ylabel='total_bill'>
```



```
In [30]: # CI ---> for removing error bars  
sns.pointplot(data=tips, x='sex', y='total_bill' ,  
              hue='smoker', ci=None)
```

```
Out[30]: <AxesSubplot:xlabel='sex', ylabel='total_bill'>
```

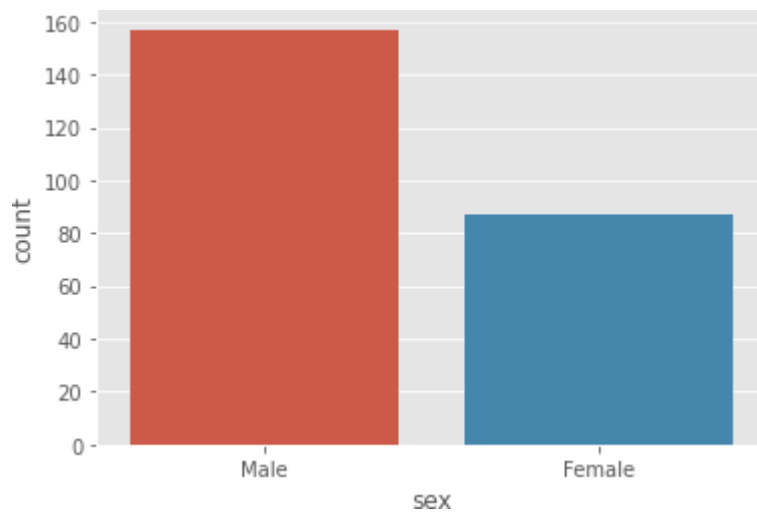


3.Count Plot

A special case for the bar plot is when you want to show the number of observations in each category rather than computing a statistic for a second variable. This is similar to a histogram over a categorical, rather than quantitative, variable

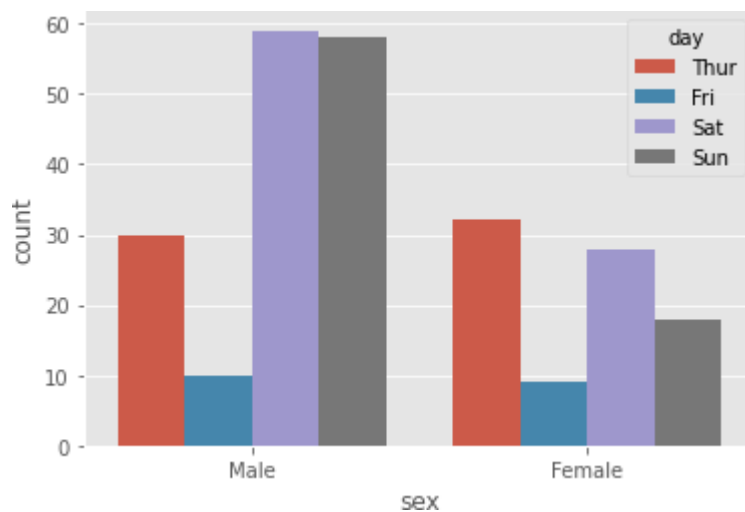
```
In [31]: # countplot  
  
sns.countplot(data=tips,x='sex')
```

Out[31]: <AxesSubplot:xlabel='sex', ylabel='count'>



```
In [32]: sns.countplot(data=tips,x='sex',hue='day')
```

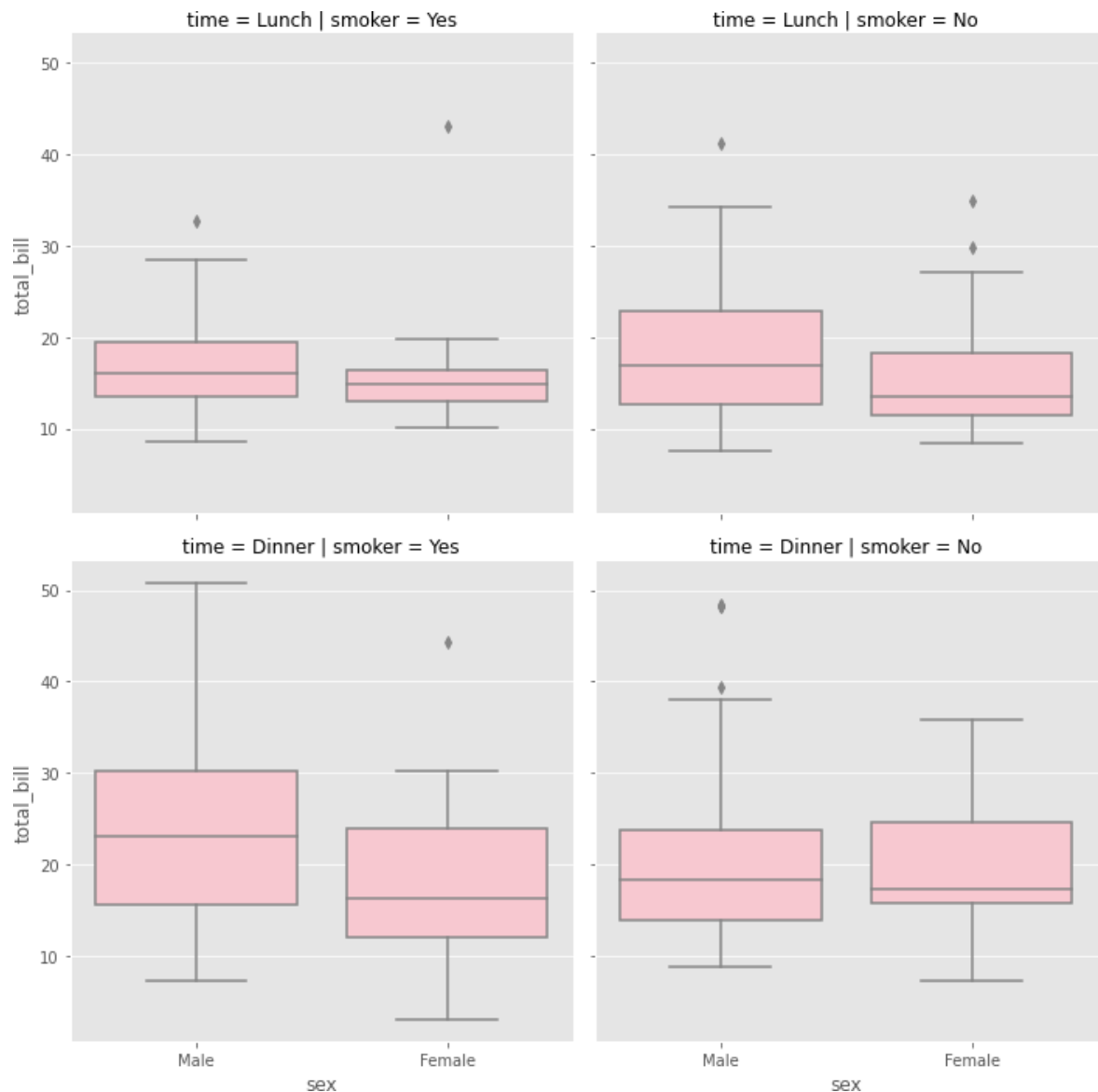
Out[32]: <AxesSubplot:xlabel='sex', ylabel='count'>



In [33]: `# faceting using catplot`

```
sns.catplot(data=tips, x='sex', y='total_bill',
            col='smoker', kind='box', row='time', color='pink')
```

Out[33]: `<seaborn.axisgrid.FacetGrid at 0x1e860f46a00>`



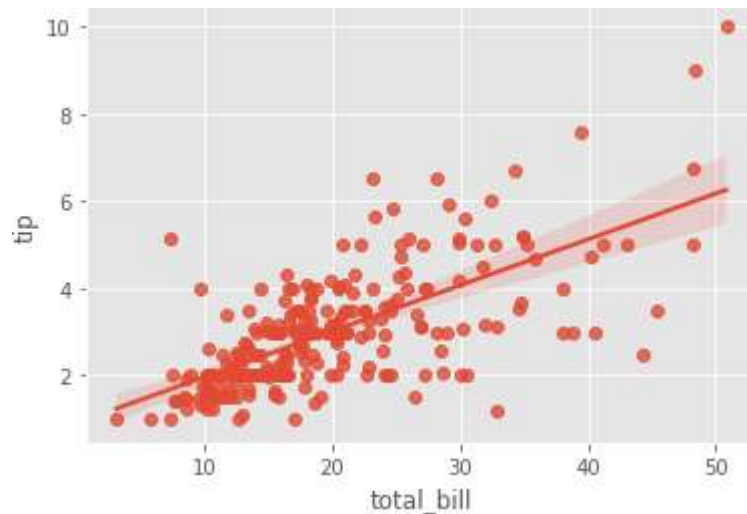
Regression Plots

- regplot
- Implot

In the simplest invocation, both functions draw a scatterplot of two variables, x and y, and then fit the regression model $y \sim x$ and plot the resulting regression line and a 95% confidence interval for that regression.

```
In [34]: # axes level  
  
# hue parameter is not available  
  
sns.regplot(data=tips,x='total_bill',y='tip')
```

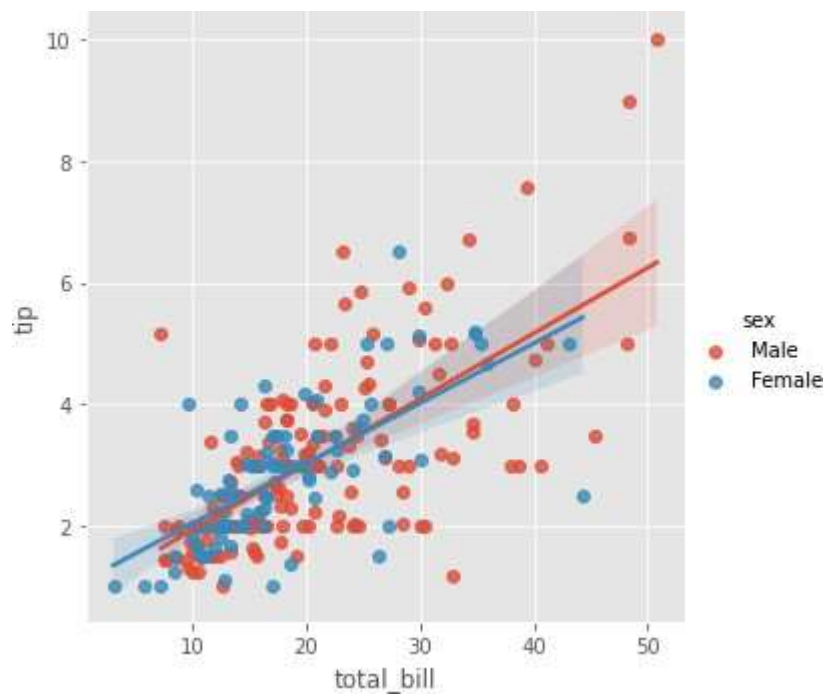
Out[34]: <AxesSubplot:xlabel='total_bill', ylabel='tip'>



Lmplot

```
In [35]: sns.lmplot(data=tips,x='total_bill',y='tip', hue = 'sex')
```

Out[35]: <seaborn.axisgrid.FacetGrid at 0x1e862649a00>

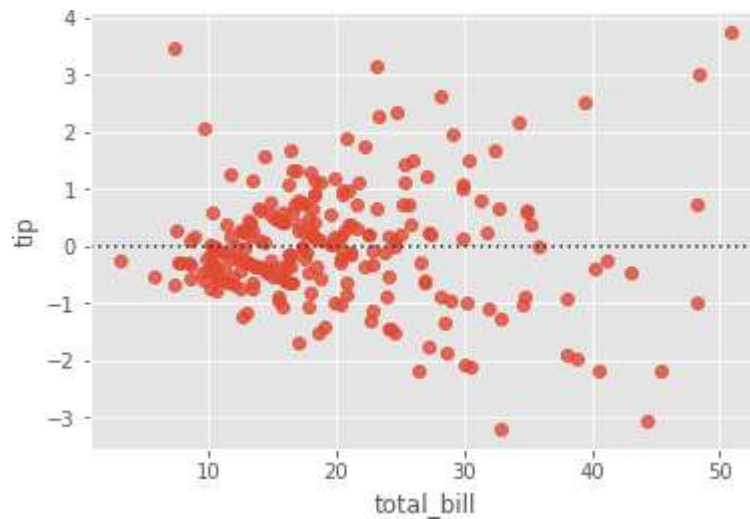


residplot

In [36]: *# residual plot*

```
sns.residplot(data=tips,x='total_bill',y='tip')
```

Out[36]: <AxesSubplot:xlabel='total_bill', ylabel='tip'>

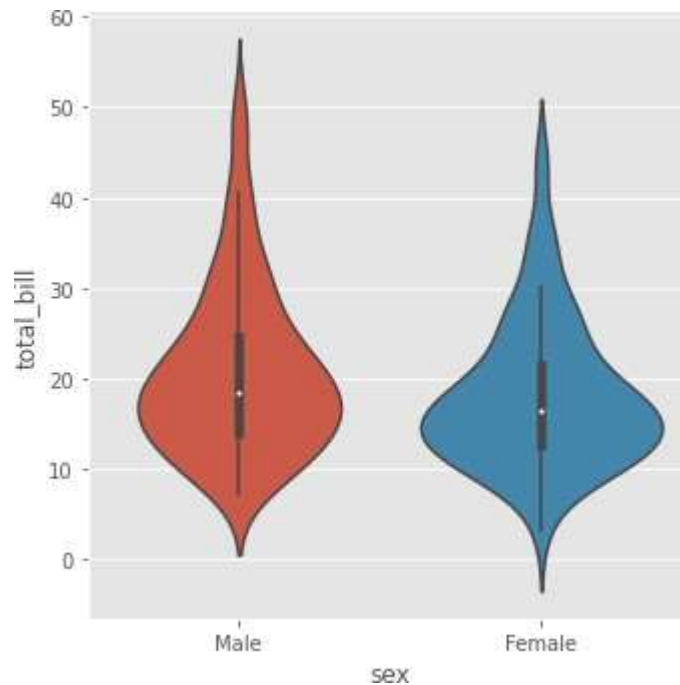


A second way to plot Facet plots -> FacetGrid

```
In [37]: # figure level -> relplot -> displot -> catplot -> lmplot

sns.catplot(data=tips,x='sex',y='total_bill',
            kind='violin')
```

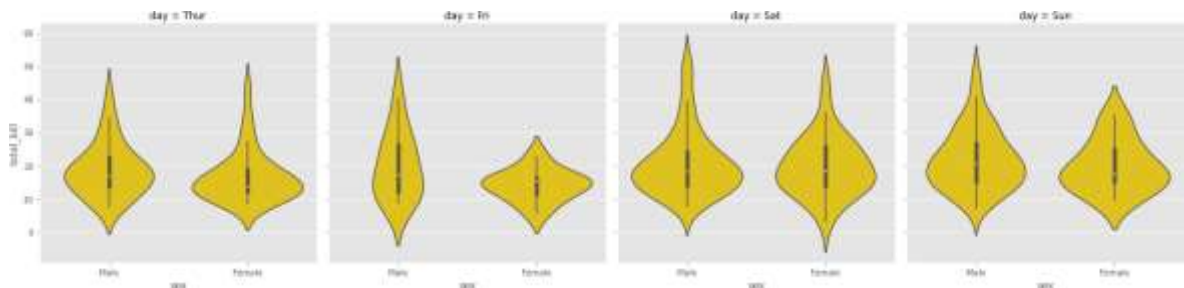
Out[37]: <seaborn.axisgrid.FacetGrid at 0x1e85e854190>



```
In [41]: # Facet Plot - columns

sns.catplot(data=tips,x='sex',y='total_bill', color = 'gold',
            kind='violin',col='day')
```

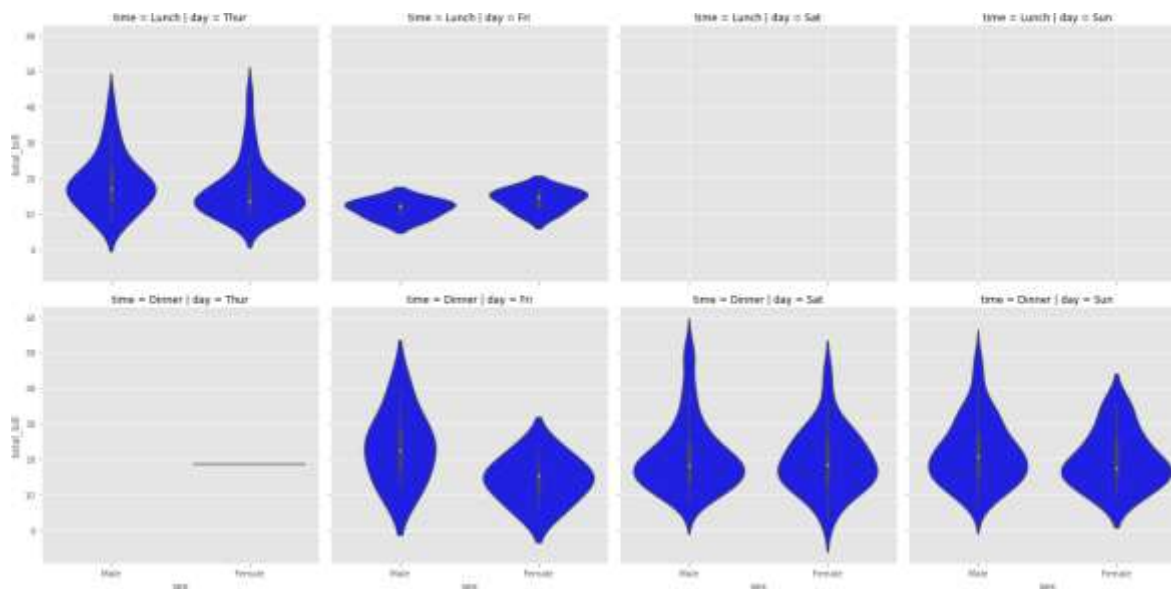
Out[41]: <seaborn.axisgrid.FacetGrid at 0x1e862e98fd0>



```
In [50]: # Facet Plot - column + rows
```

```
sns.catplot(data=tips,x='sex',y='total_bill',color='blue',  
            kind='violin',col='day',row='time')
```

```
Out[50]: <seaborn.axisgrid.FacetGrid at 0x1e86965dbe0>
```



Facetgrid

```
In [48]: # Second Method - Facetgrid

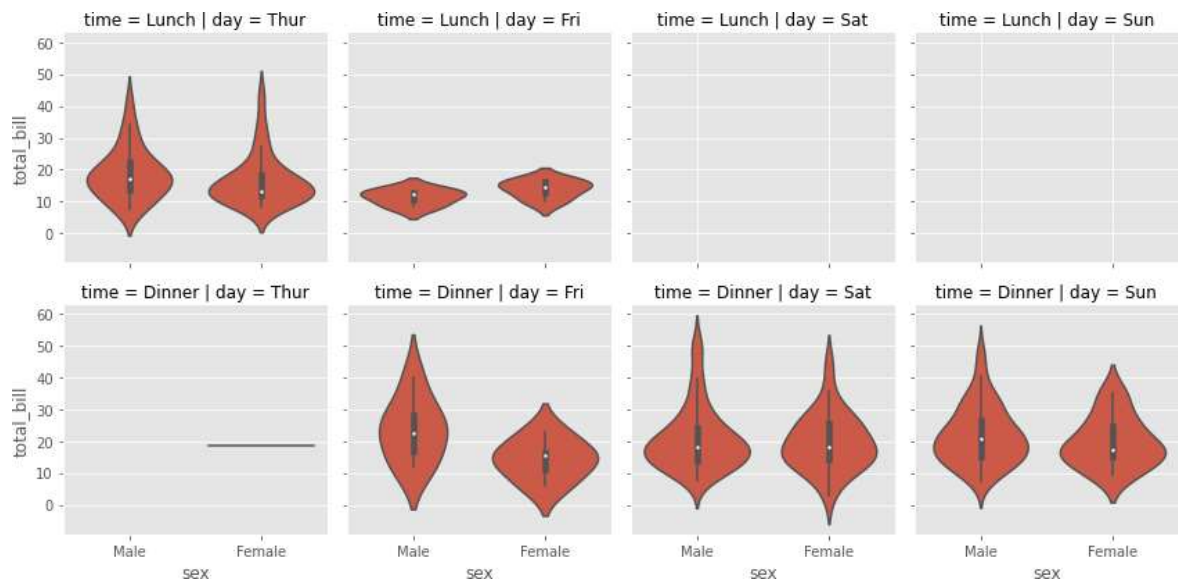
# it is the Lower Level code of catplot

secondmethod = sns.FacetGrid(data =tips ,col ='day' , row ='time')

secondmethod.map(sns.violinplot,'sex','total_bill')
```

C:\Users\user\anaconda3\lib\site-packages\seaborn\axisgrid.py:670: UserWarning: Using the violinplot function without specifying `order` is likely to produce an incorrect plot.
warnings.warn(warning)

Out[48]: <seaborn.axisgrid.FacetGrid at 0x1e868e51580>

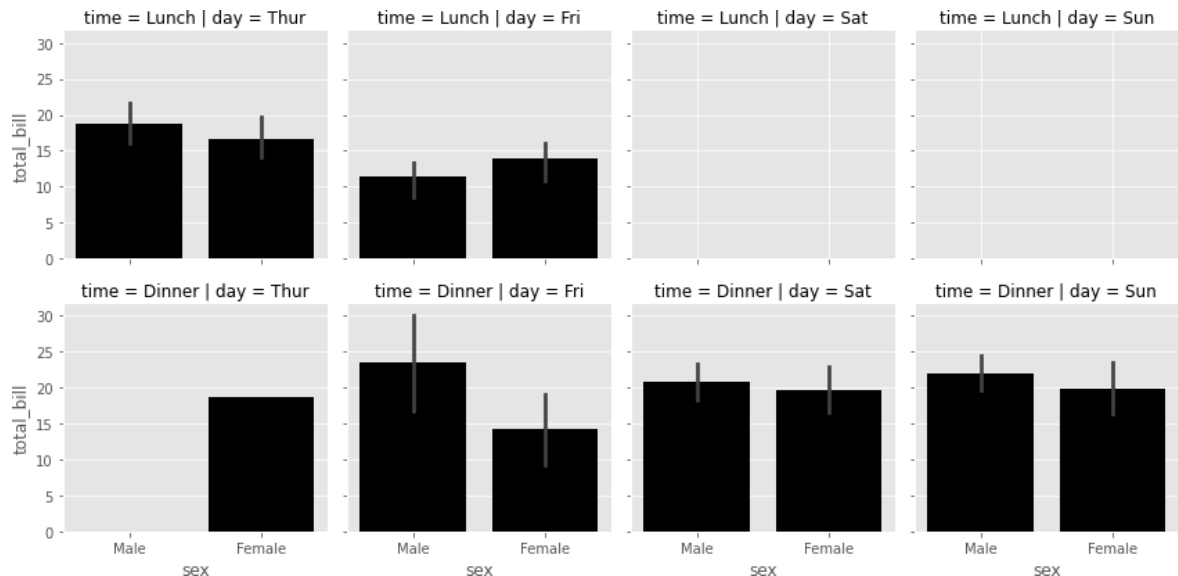



```
In [53]: secondmethod = sns.FacetGrid(data =tips ,col = 'day' , row = 'time' )

secondmethod.map(sns.barplot, 'sex', 'total_bill', color='black')
```

C:\Users\user\anaconda3\lib\site-packages\seaborn\axisgrid.py:670: UserWarning: Using the barplot function without specifying `order` is likely to produce an incorrect plot.
warnings.warn(warning)

Out[53]: <seaborn.axisgrid.FacetGrid at 0x1e867b9caf0>



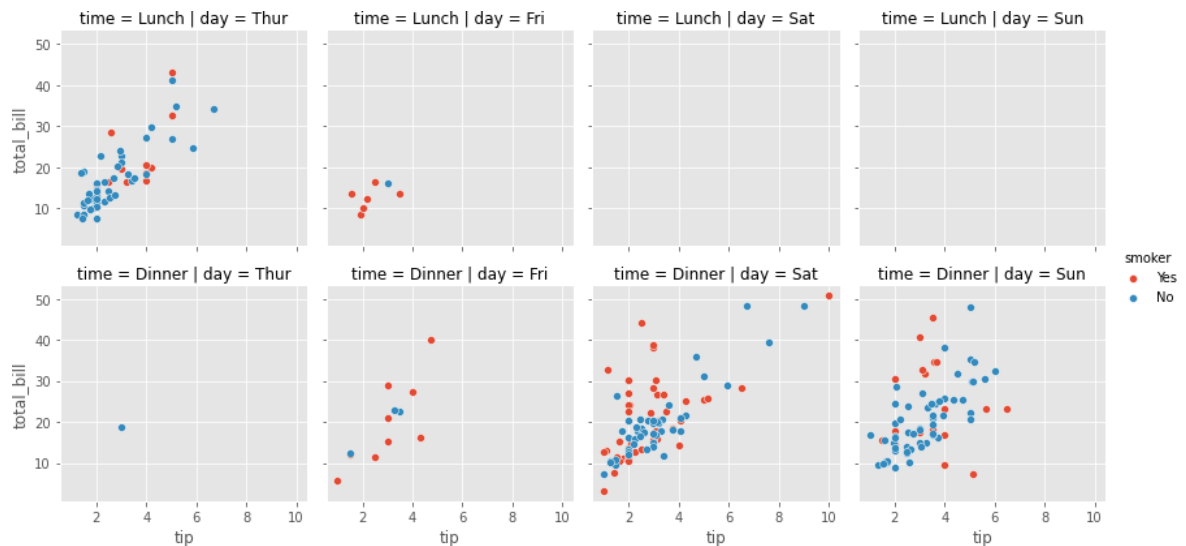
```
In [57]: # Adding Legend

secondmethod = sns.FacetGrid(data=tips,col='day',row='time',hue='smoker')

secondmethod.map(sns.scatterplot, 'tip', 'total_bill')

secondmethod.add_legend()
```

Out[57]: <seaborn.axisgrid.FacetGrid at 0x1e86bf707f0>



Plotting Pairwise Relationship (PairGrid Vs Pairplot)

Here's a concise explanation:

- **Pairplot:** It is a function in the seaborn library that **creates a grid of scatterplots**, showing the relationships between variables in a dataset. Each variable is plotted against every other variable, providing a visual overview of their correlations.
- **PairGrid:** It is a class in seaborn that allows for **more customization** in creating a grid of subplots. With PairGrid, you can manually specify the plots to be displayed on the grid, including scatterplots, histograms, kernel density plots, etc. It provides more flexibility in creating complex visualizations compared to pairplot.

In [59]: `iris.sample()`

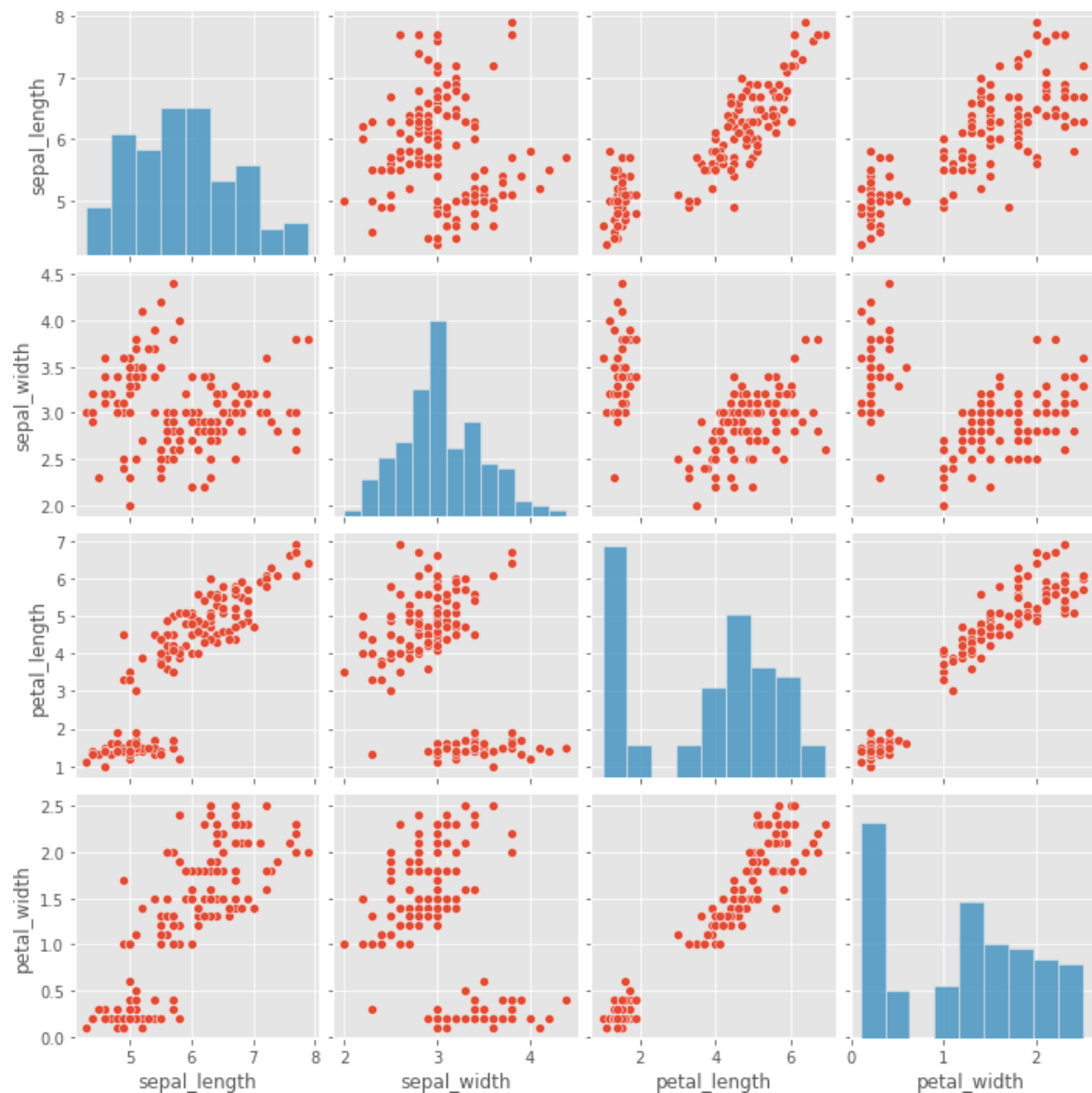
Out[59]:

| | sepal_length | sepal_width | petal_length | petal_width | species |
|----|--------------|-------------|--------------|-------------|------------|
| 53 | 5.5 | 2.3 | 4.0 | 1.3 | versicolor |

```
In [60]: # Pairplot
```

```
sns.pairplot(iris)
```

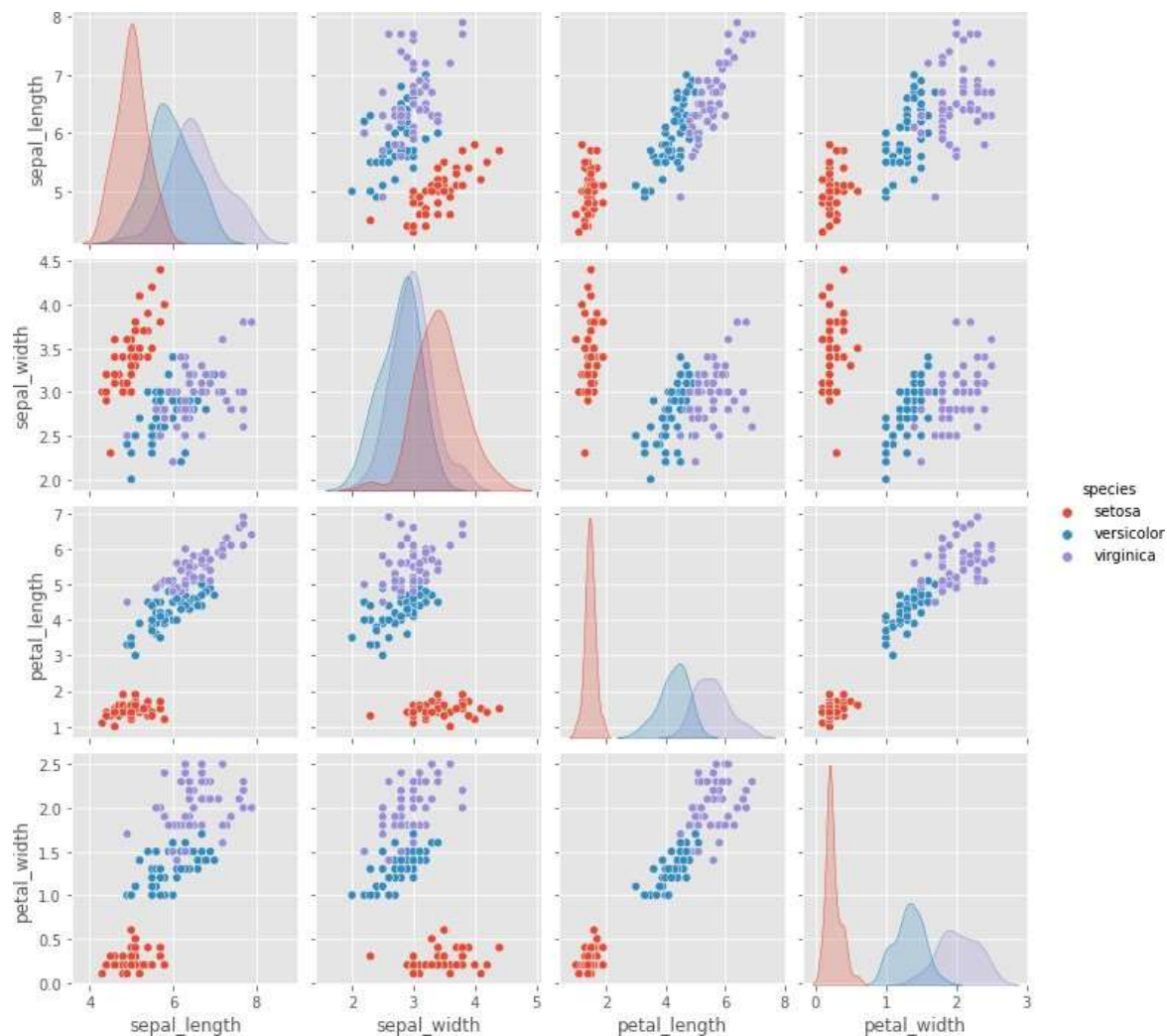
```
Out[60]: <seaborn.axisgrid.PairGrid at 0x1e86c618880>
```



```
In [61]: # hue
```

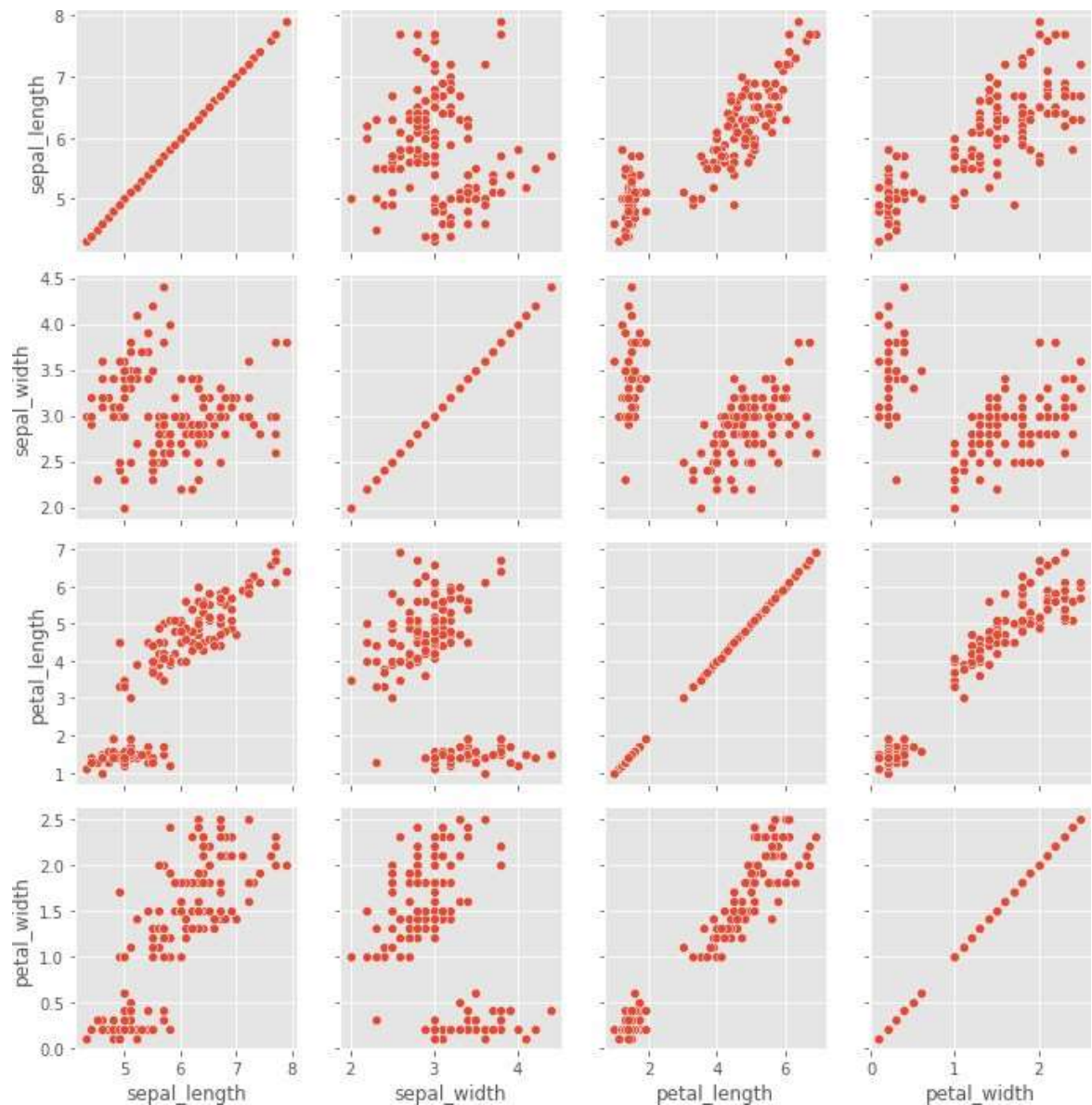
```
sns.pairplot(iris,hue='species')
```

```
Out[61]: <seaborn.axisgrid.PairGrid at 0x1e86da65460>
```



```
In [63]: # pair grid  
  
g = sns.PairGrid(data=iris )  
  
# g.map  
  
g.map(sns.scatterplot)
```

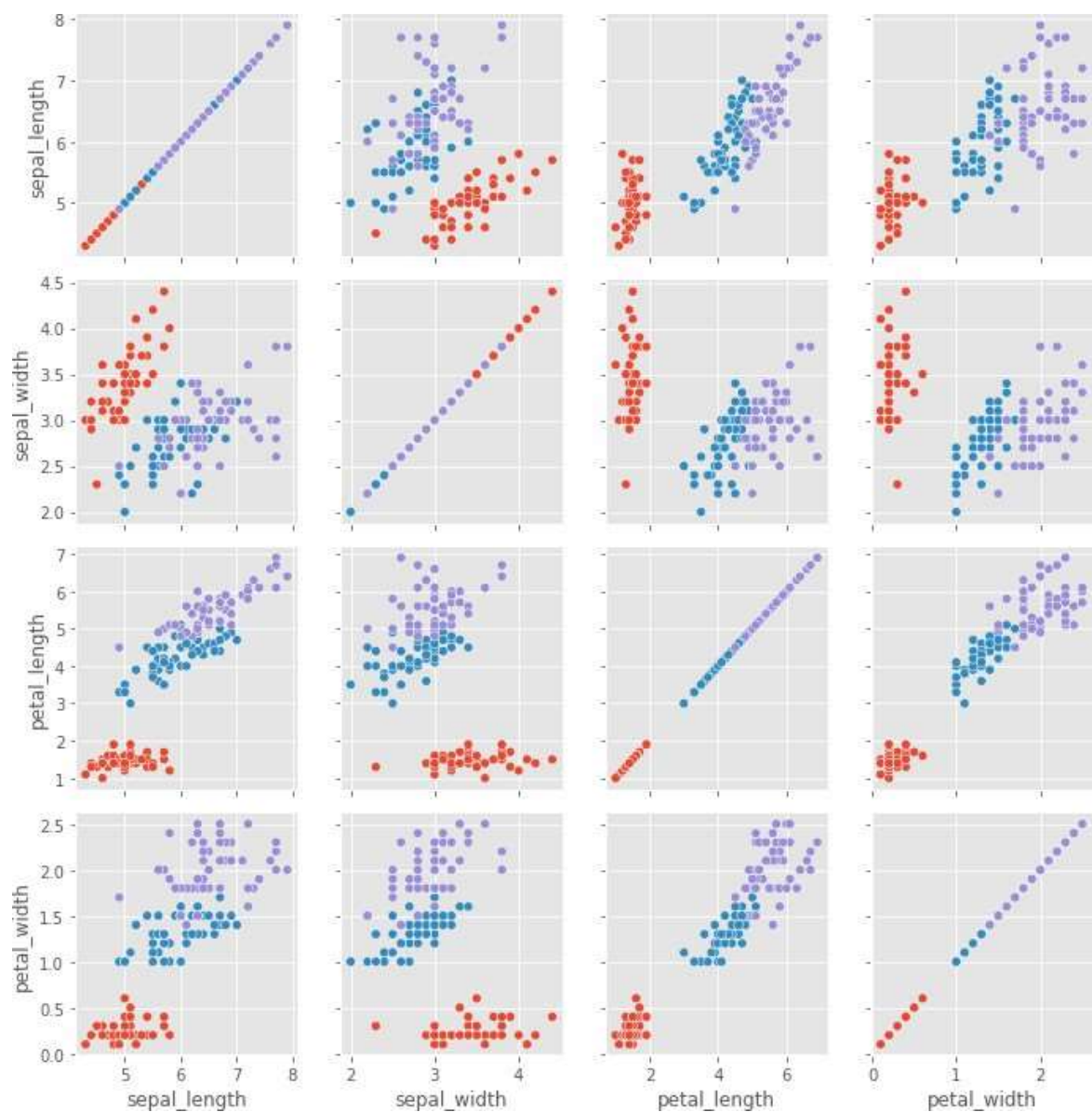
Out[63]: <seaborn.axisgrid.PairGrid at 0x1e8707bed90>



```
In [64]: # hue
g = sns.PairGrid(data=iris,hue='species')

# g.map
g.map(sns.scatterplot)
```

Out[64]: <seaborn.axisgrid.PairGrid at 0x1e870873640>

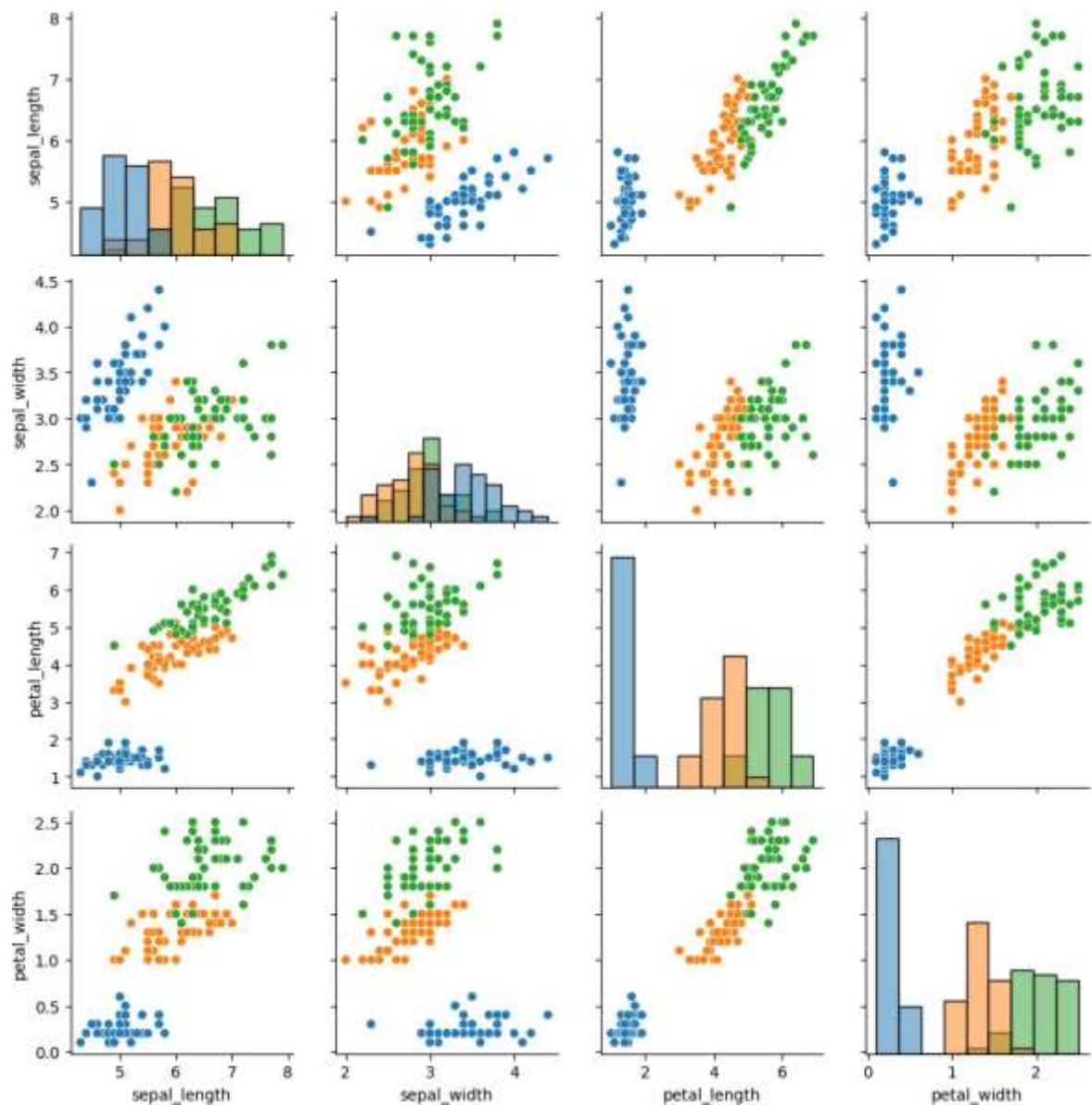


```
In [66]: # style
plt.style.use('default')
```



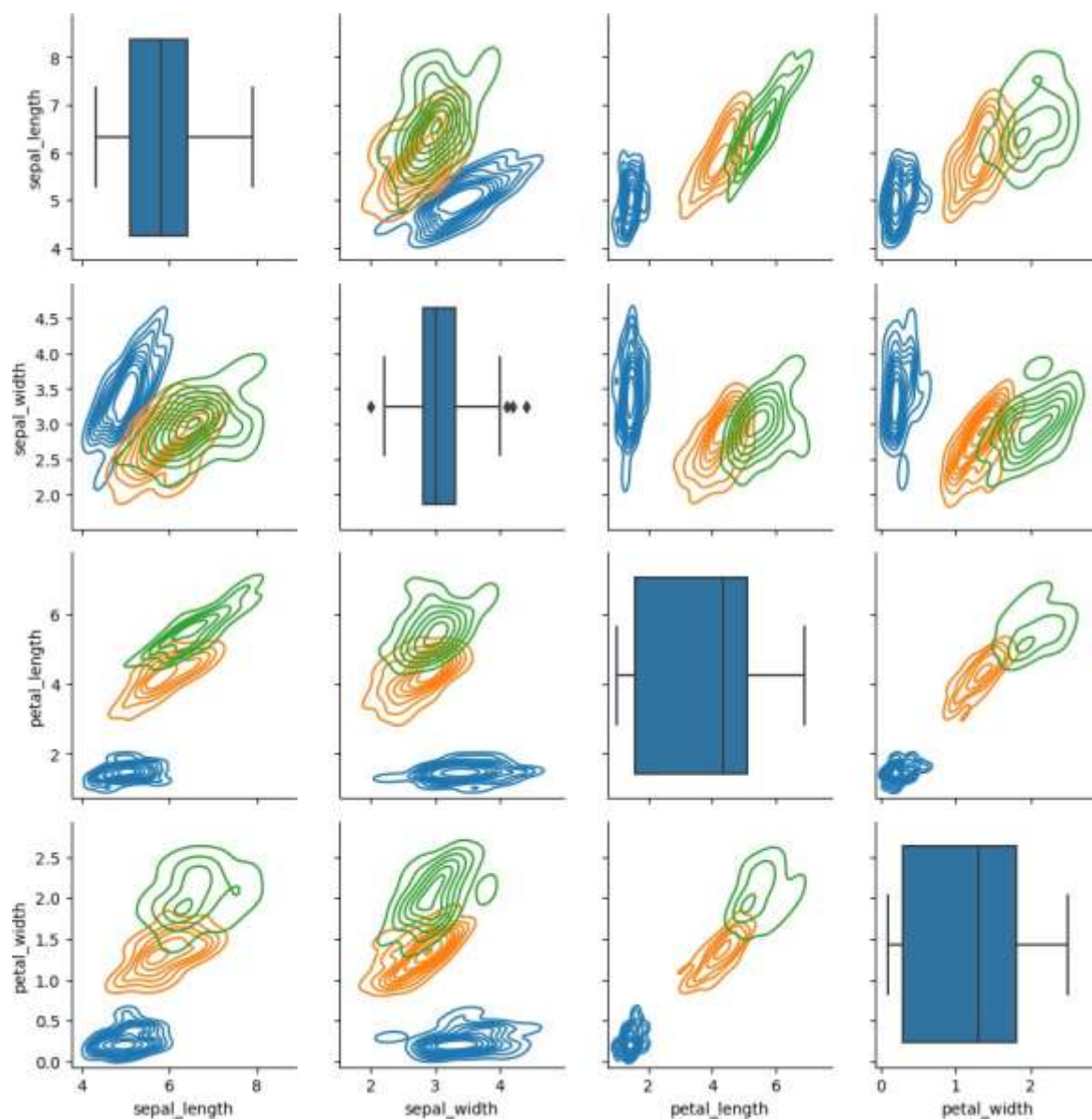
```
In [68]: # map_diag -> map_offdiag  
g = sns.PairGrid(data=iris,hue='species')  
g.map_diag(sns.histplot)  
g.map_offdiag(sns.scatterplot)
```

Out[68]: <seaborn.axisgrid.PairGrid at 0x1e8759bbc10>



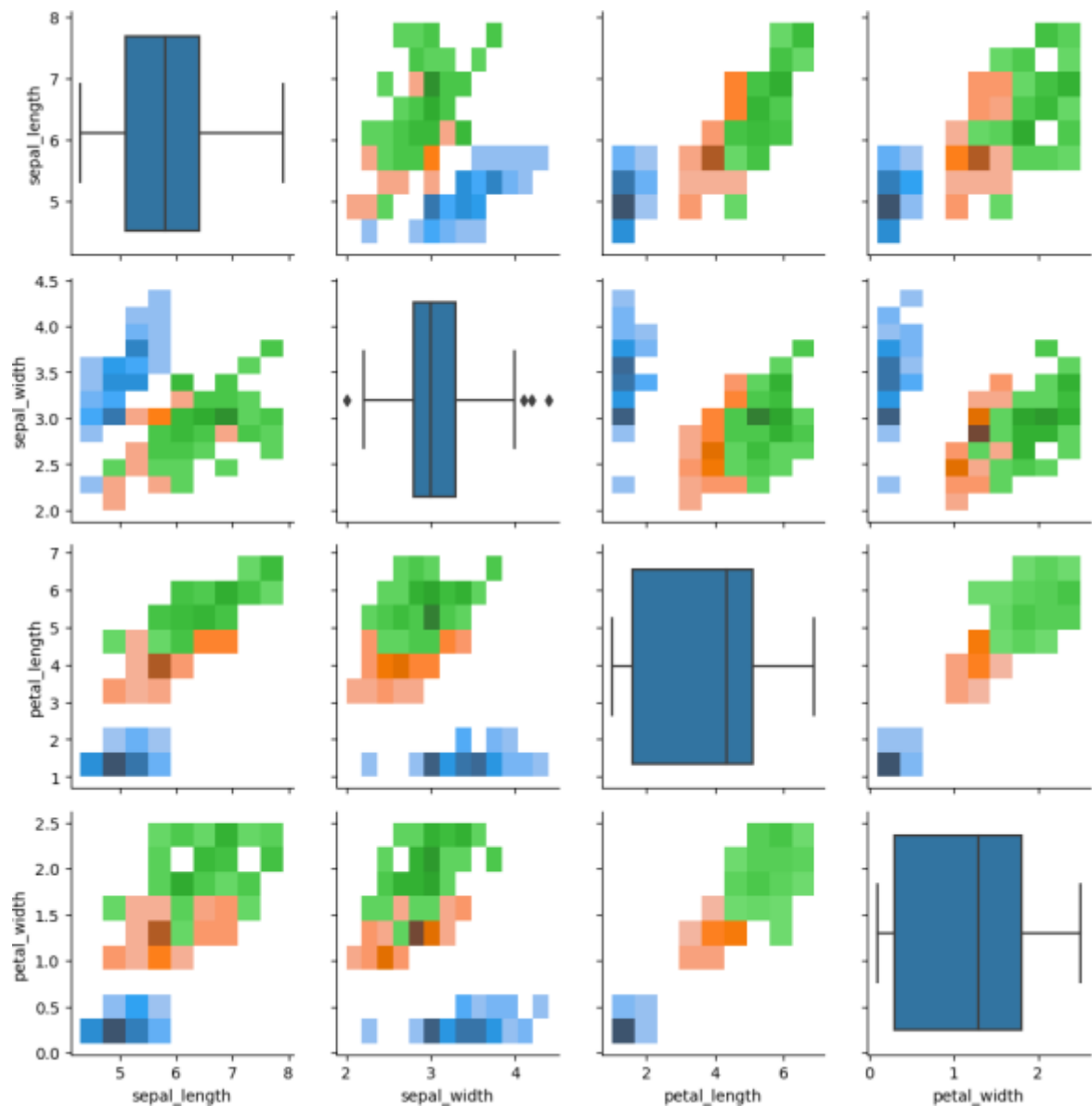
```
In [69]: g = sns.PairGrid(data=iris,hue='species')  
g.map_diag(sns.boxplot)  
g.map_offdiag(sns.kdeplot)
```

Out[69]: <seaborn.axisgrid.PairGrid at 0x1e87684fca0>



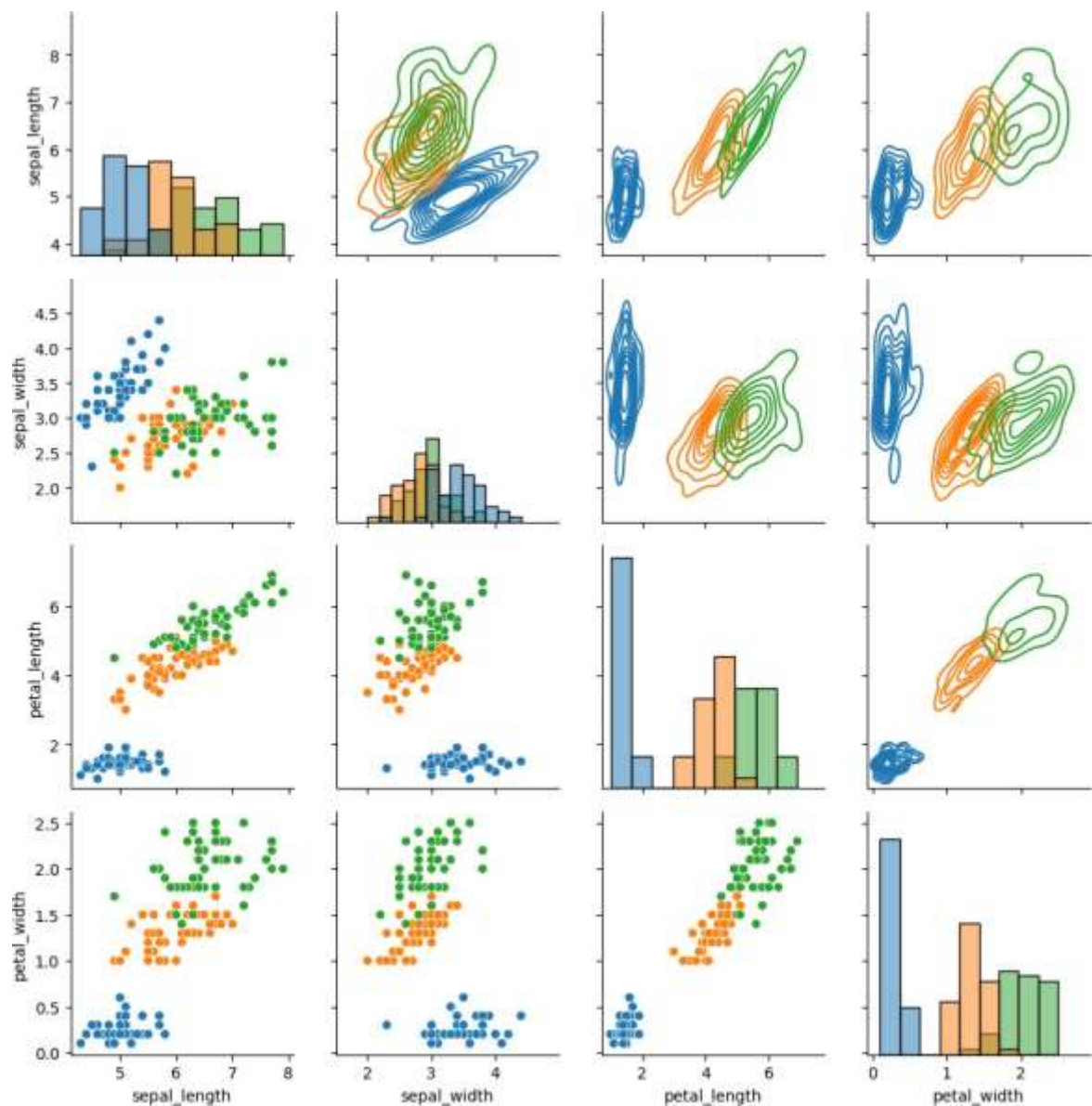

```
In [71]: g = sns.PairGrid(data=iris,hue='species')
g.map_diag(sns.boxplot)
g.map_offdiag(sns.histplot)
```

Out[71]: <seaborn.axisgrid.PairGrid at 0x1e8788cdb20>



```
In [72]: # map_diag -> map_upper -> map_lower
g = sns.PairGrid(data=iris,hue='species')
g.map_diag(sns.histplot) # diagonal
g.map_upper(sns.kdeplot) # upper
g.map_lower(sns.scatterplot) # lower
```

Out[72]: <seaborn.axisgrid.PairGrid at 0x1e8788de760>



```
In [73]: # vars - working upon the desired columns

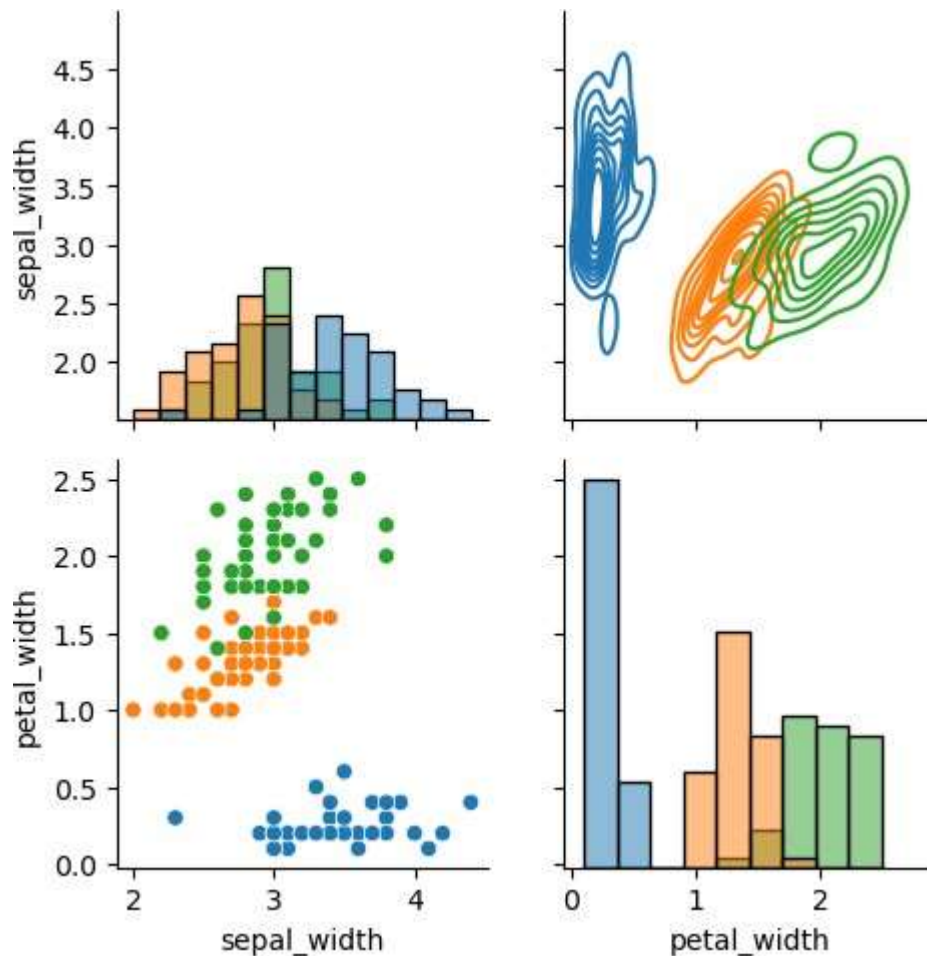
g = sns.PairGrid(data=iris,hue='species',vars=['sepal_width','petal_width'])

g.map_diag(sns.histplot)

g.map_upper(sns.kdeplot)

g.map_lower(sns.scatterplot)
```

Out[73]: <seaborn.axisgrid.PairGrid at 0x1e87b757520>



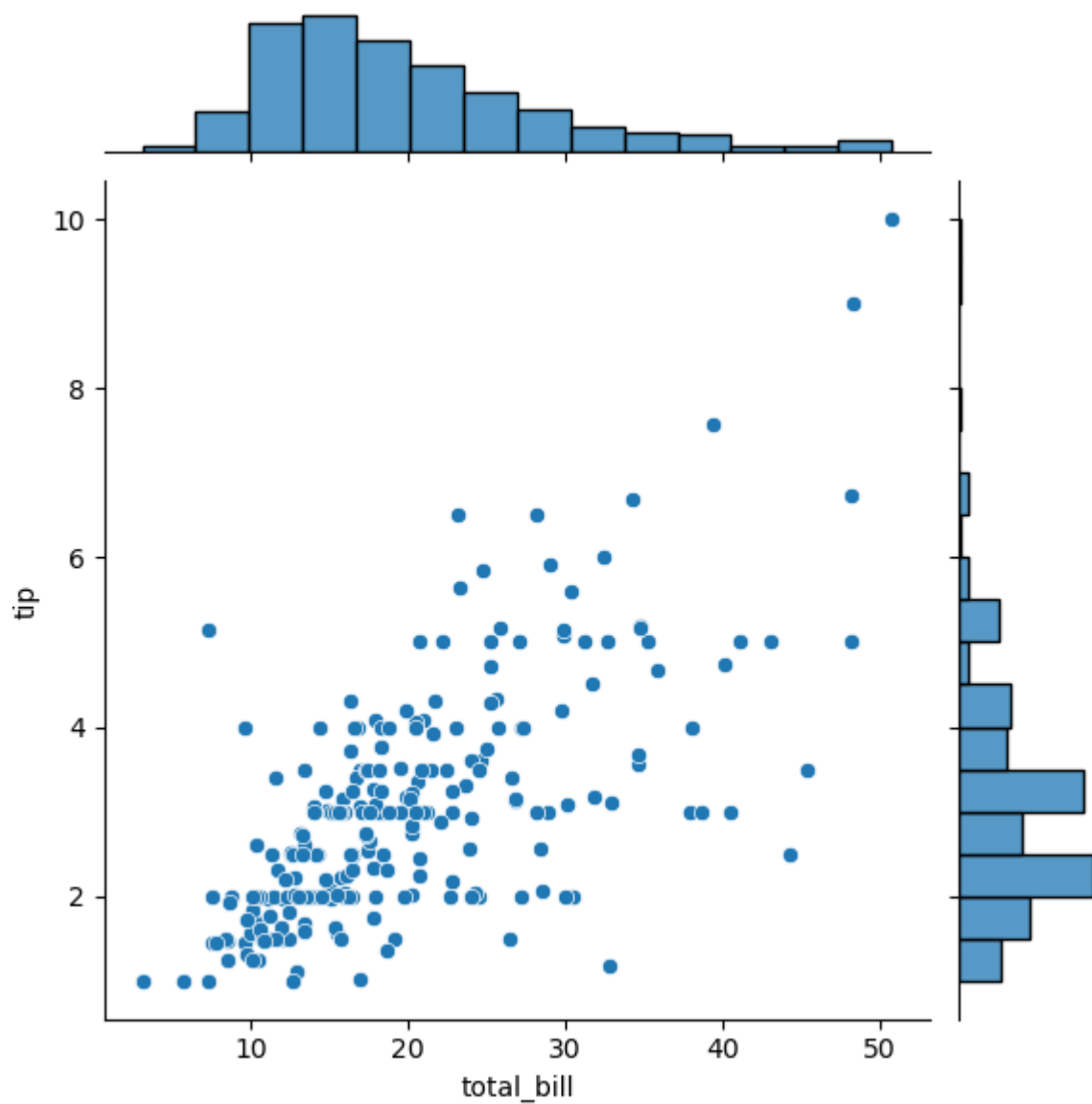
JointGrid Vs Jointplot

Here's a concise explanation:

- **Jointplot** : It is a function in seaborn that combines scatterplot and histograms (or kernel density plots) to visualize the **relationship between two variables**. It shows both the individual distributions and the correlation between the variables.
- **JointGrid**: It is a class in seaborn that provides a framework for creating **custom joint plots**. With JointGrid, you have more control over the layout and types of plots used in a

```
In [74]: sns.jointplot(data=tips,x='total_bill',y='tip')
```

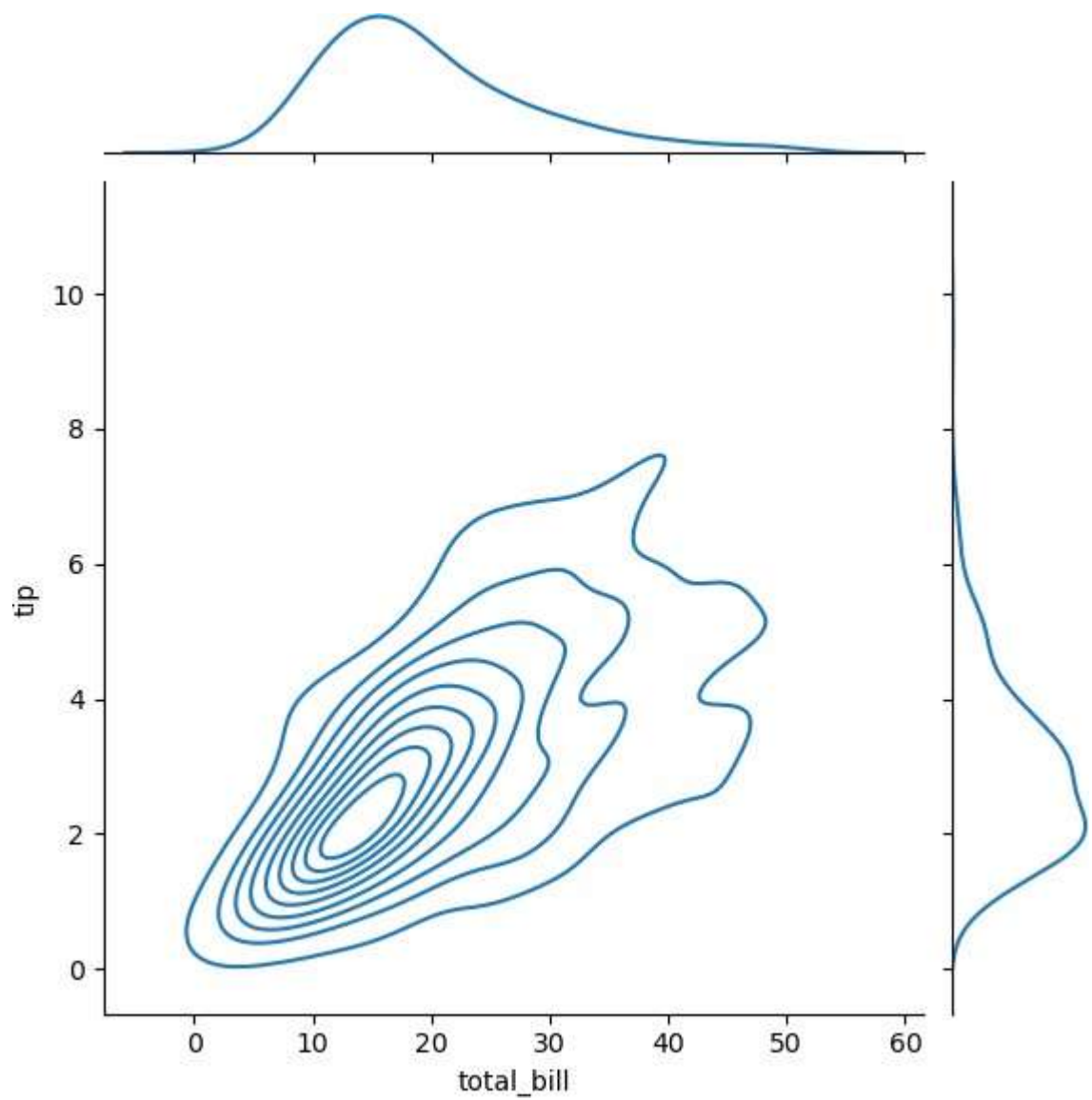
```
Out[74]: <seaborn.axisgrid.JointGrid at 0x1e87b78ad30>
```



```
In [78]: # kind
```

```
sns.jointplot(data=tips,x='total_bill',y='tip',kind= 'kde' )
```

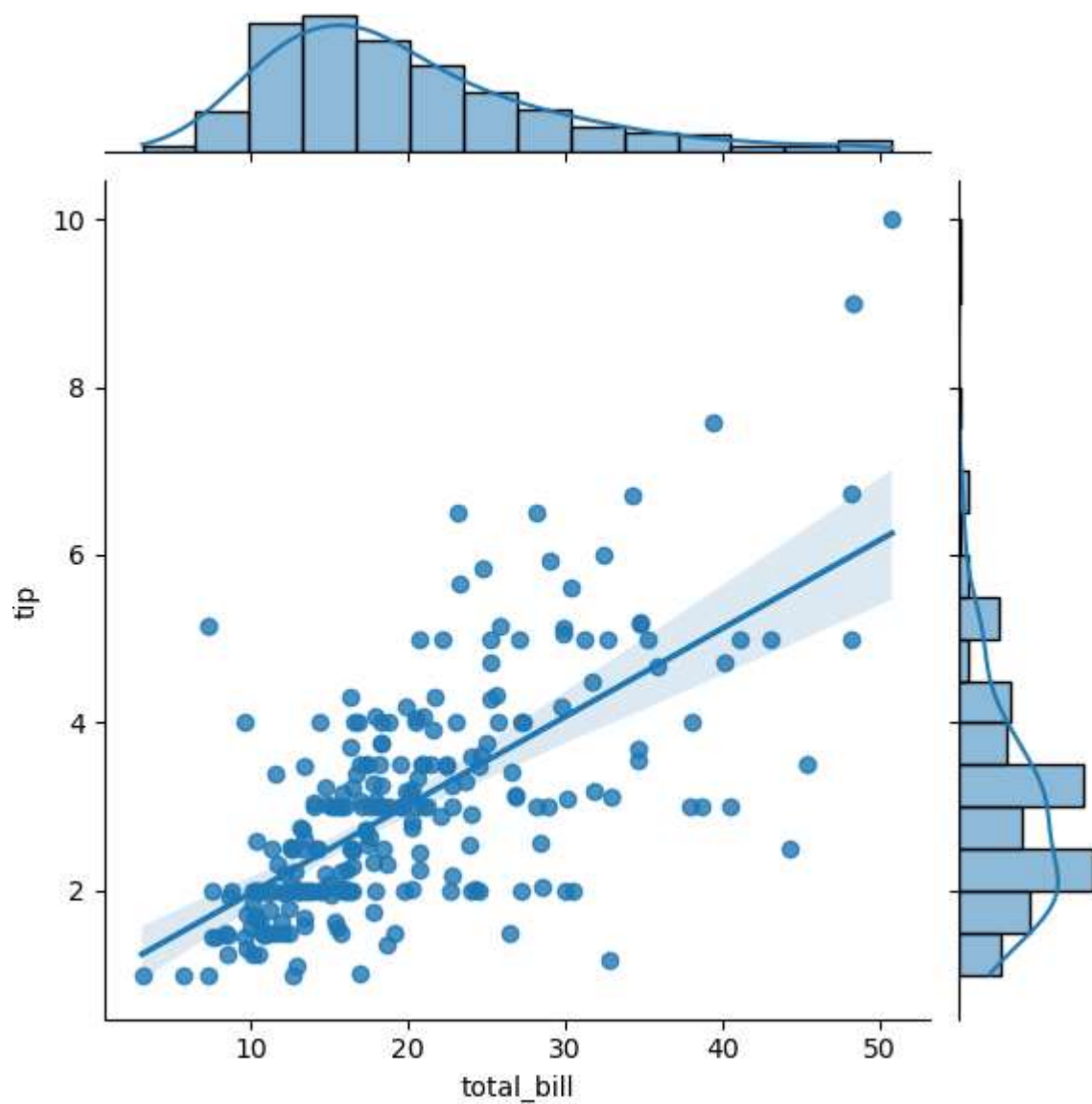
```
Out[78]: <seaborn.axisgrid.JointGrid at 0x1e8768a06a0>
```



In [83]: *#regression line*

```
sns.jointplot(data=tips,x='total_bill',y='tip',kind= 'reg' )
```

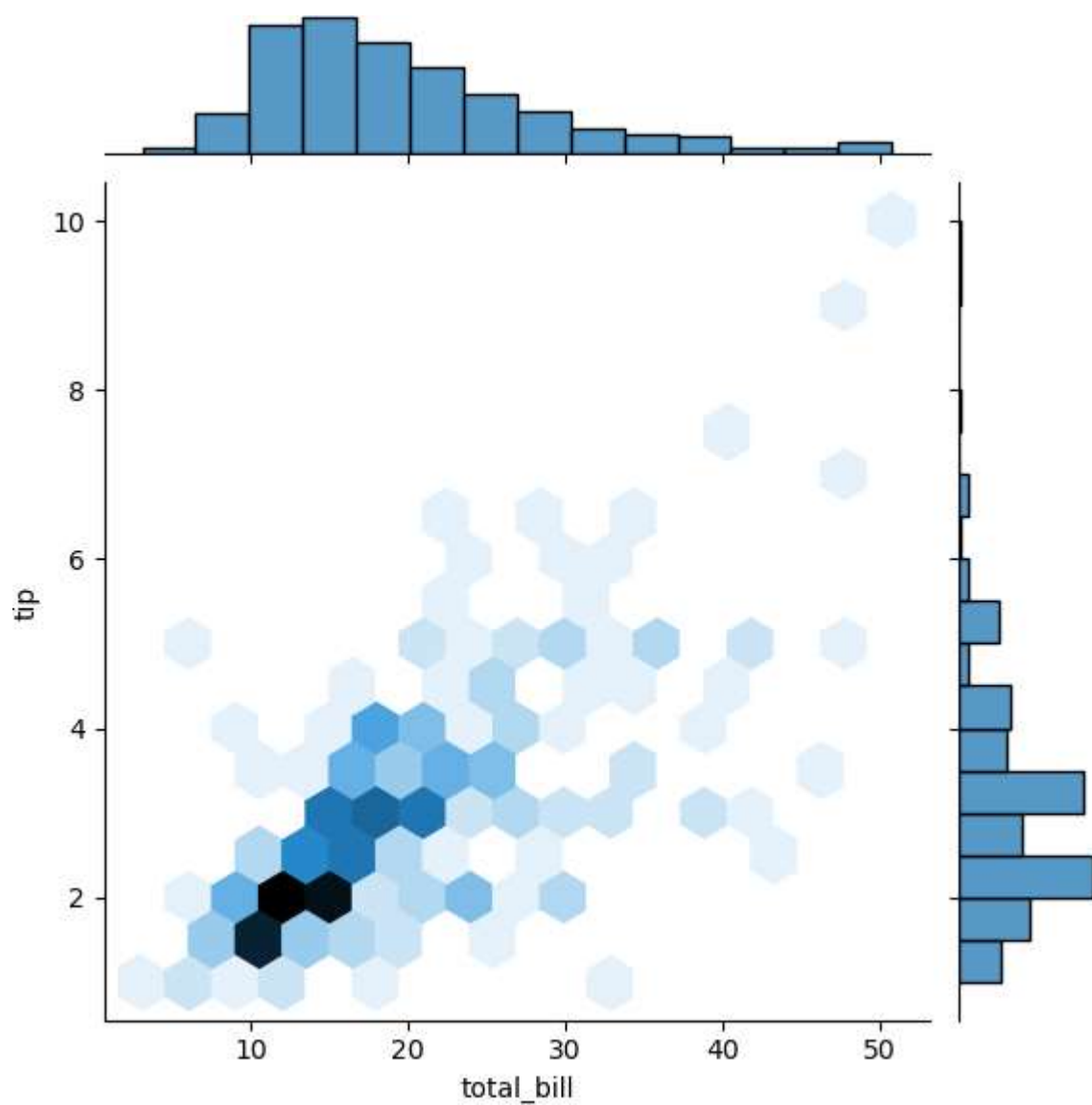
Out[83]: <seaborn.axisgrid.JointGrid at 0x1e87de2cac0>



```
In [82]: # hex
```

```
sns.jointplot(data=tips,x='total_bill',y='tip',kind= 'hex' )
```

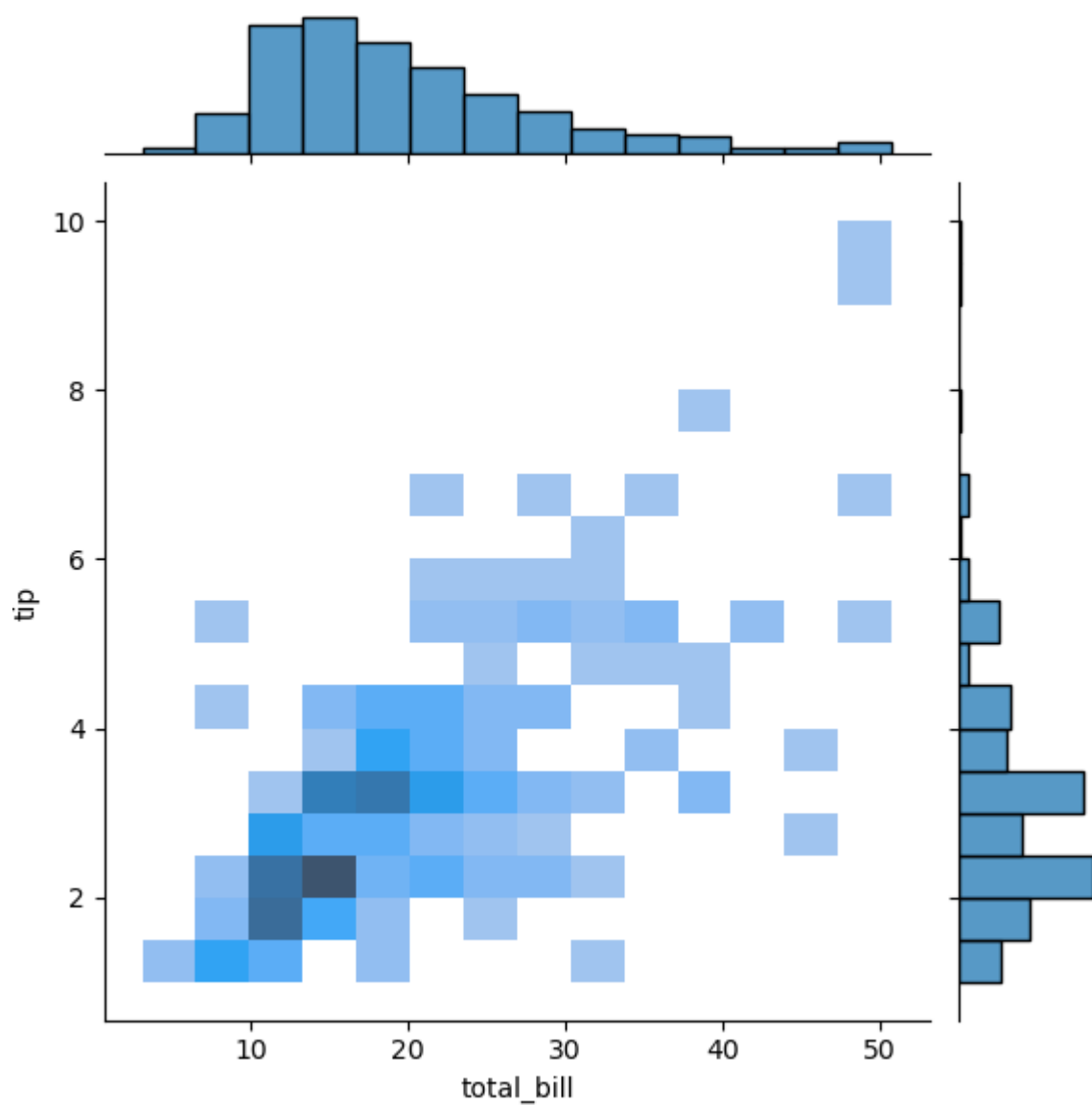
```
Out[82]: <seaborn.axisgrid.JointGrid at 0x1e87d97b370>
```



```
In [79]: # kind -2d histogram + 1 histogram
```

```
sns.jointplot(data=tips,x='total_bill',y='tip',kind='hist' )
```

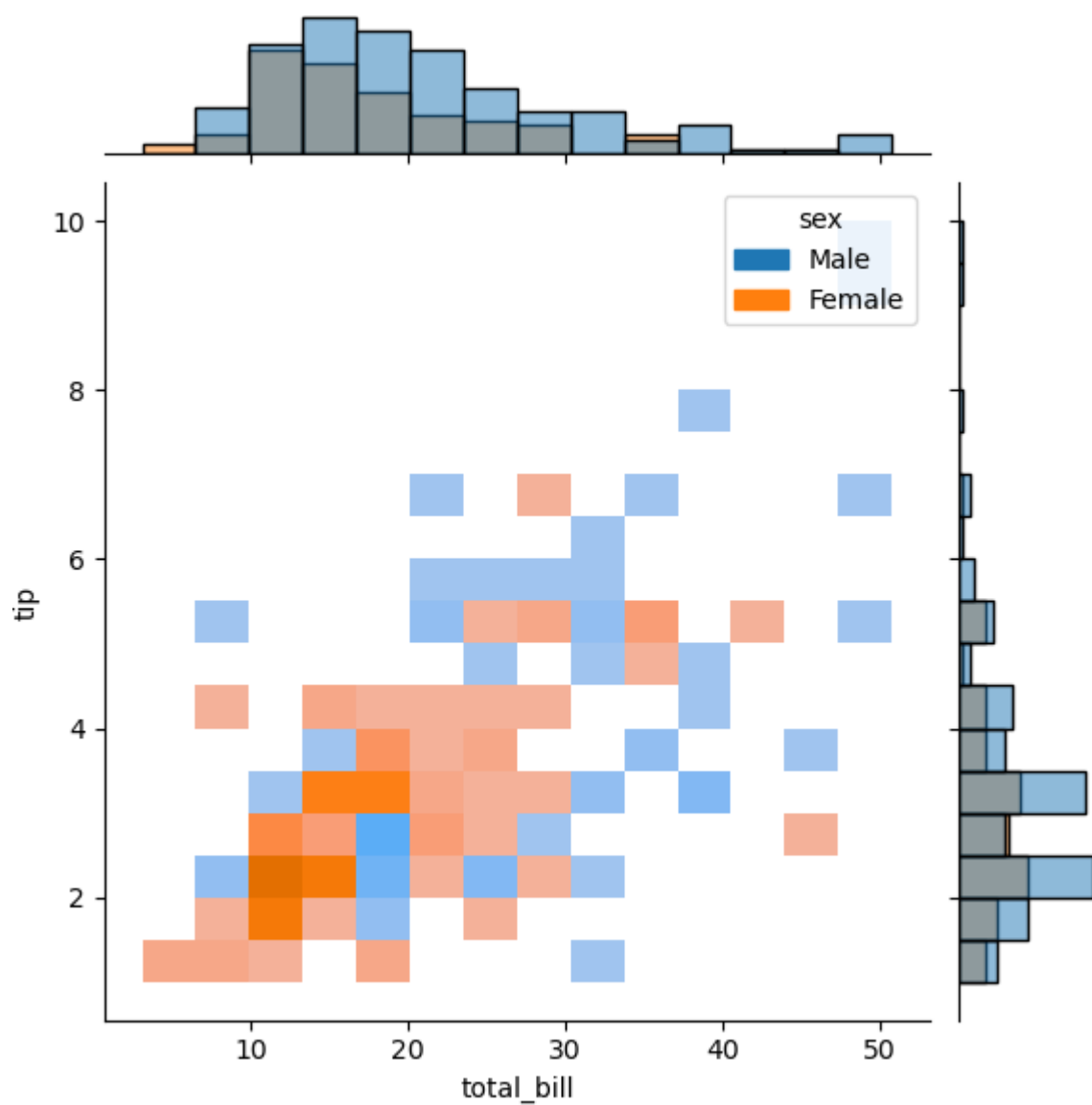
```
Out[79]: <seaborn.axisgrid.JointGrid at 0x1e877fe9250>
```




```
In [76]: # hue
```

```
sns.jointplot(data=tips,x='total_bill',y='tip',kind='hist',hue='sex')
```

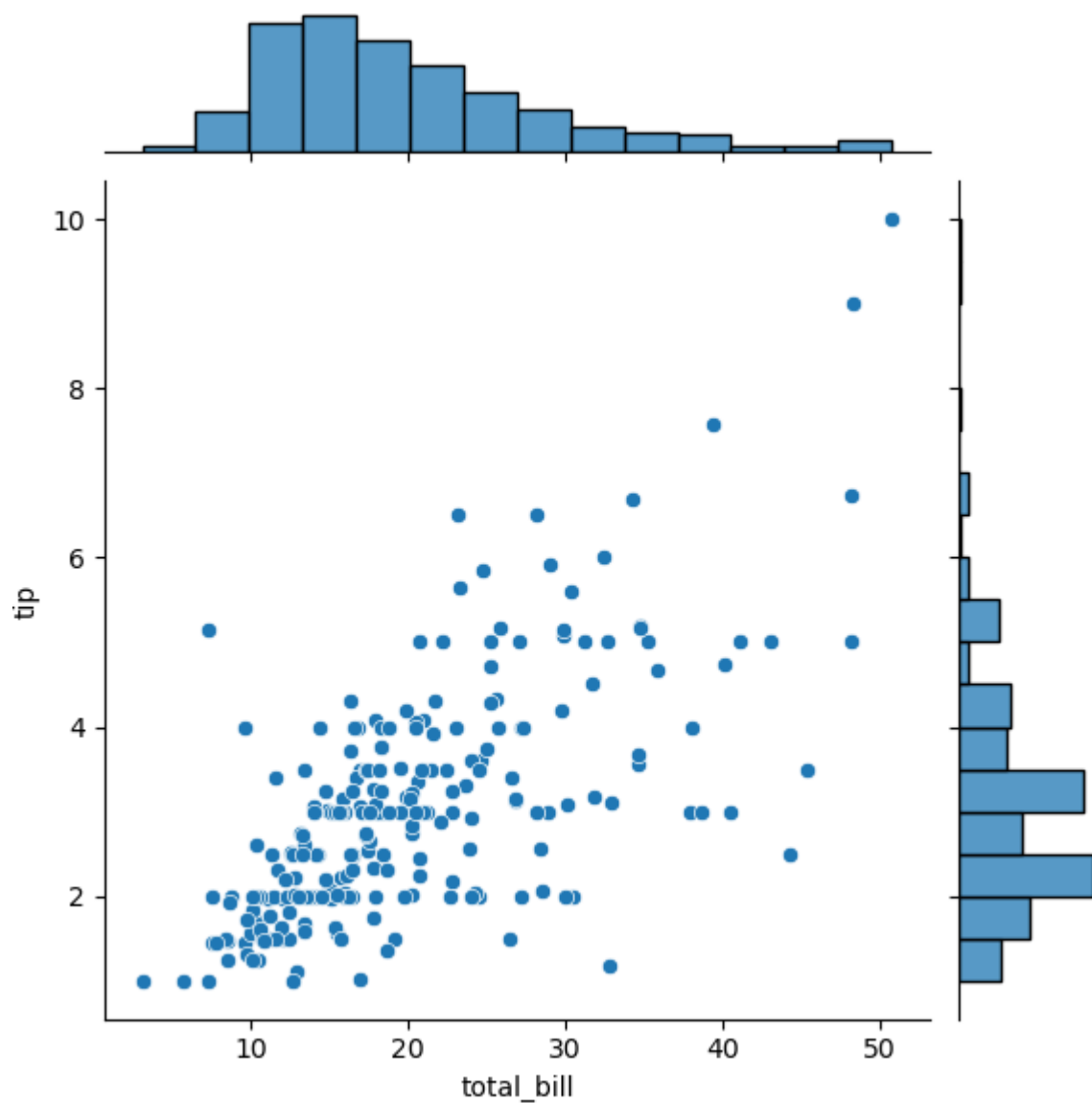
```
Out[76]: <seaborn.axisgrid.JointGrid at 0x1e878736550>
```



Jointgrid

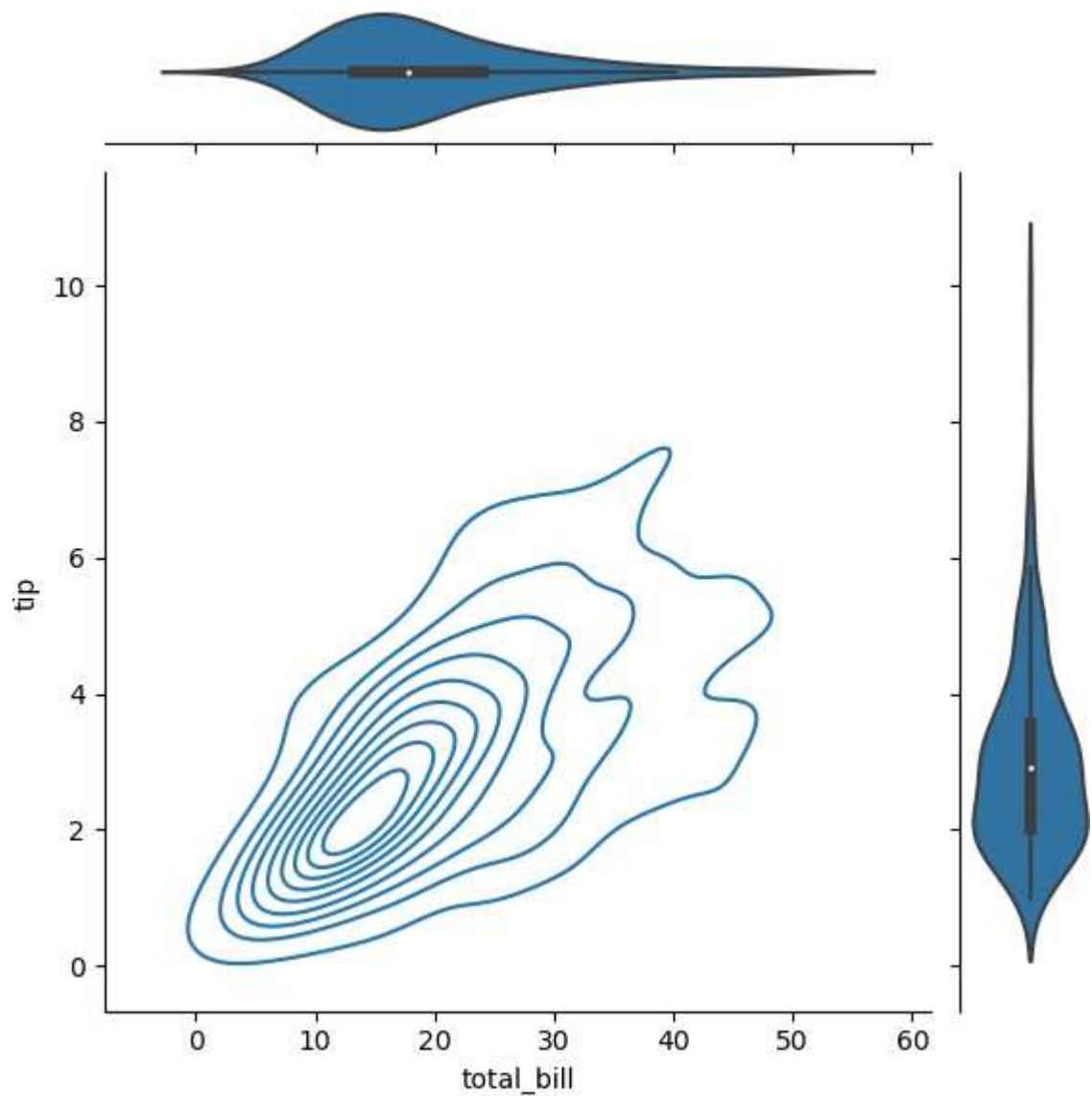
```
In [84]: # we can change plots as per our requirements  
g = sns.JointGrid(data=tips,x='total_bill',y='tip')  
g.plot(sns.scatterplot,sns.histplot)
```

```
Out[84]: <seaborn.axisgrid.JointGrid at 0x1e87e238850>
```



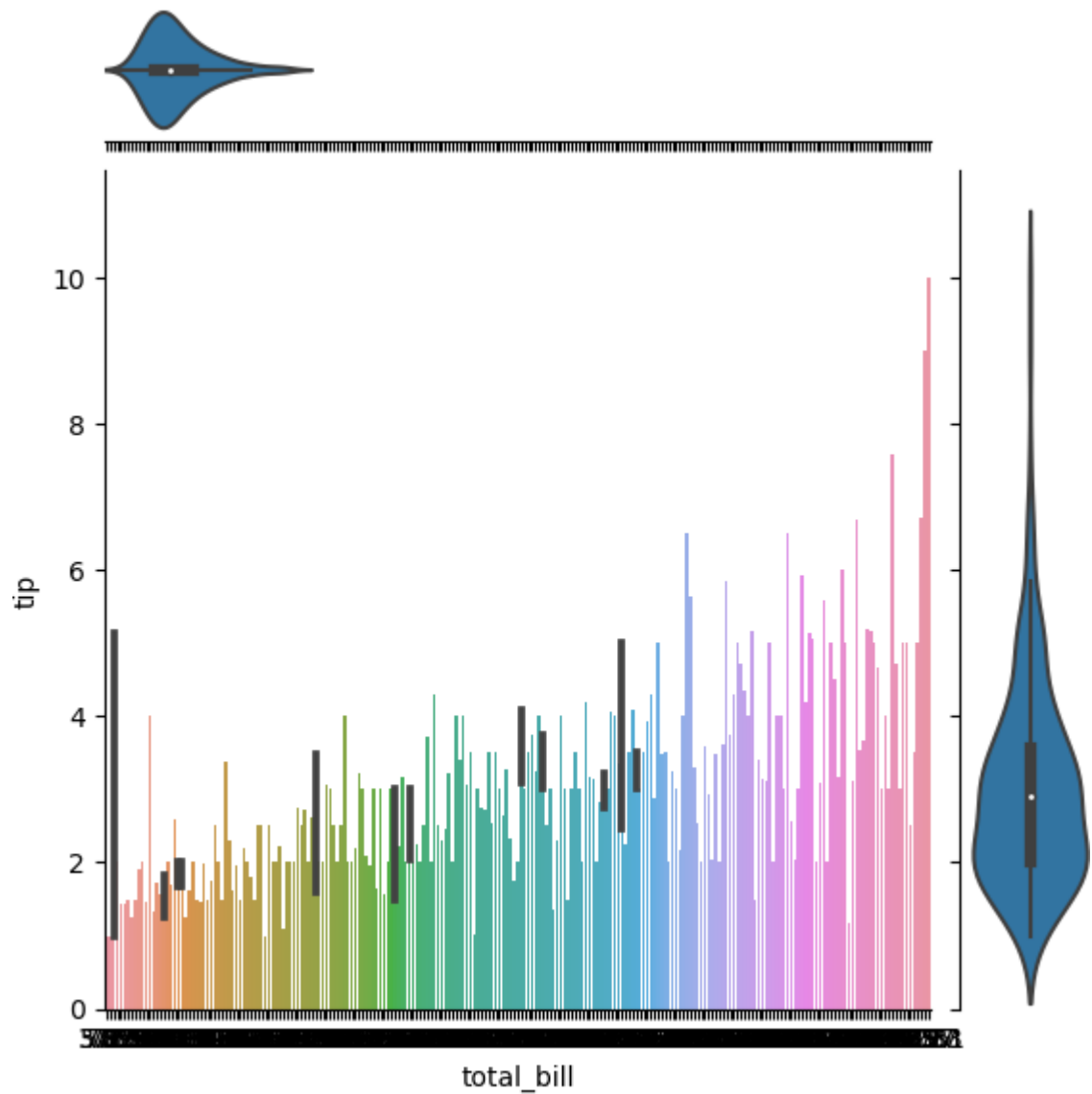
```
In [85]: g = sns.JointGrid(data=tips,x='total_bill',y='tip')  
g.plot(sns.kdeplot,sns.violinplot)
```

```
Out[85]: <seaborn.axisgrid.JointGrid at 0x1e87e7d5a90>
```



```
In [87]: g = sns.JointGrid(data=tips,x='total_bill',y='tip')  
g.plot(sns.barplot,sns.violinplot)
```

Out[87]: <seaborn.axisgrid.JointGrid at 0x1e87ec37040>



Utility Functions

In [88]: `# get dataset names`

```
sns.get_dataset_names()
```

Out[88]:

```
['anagrams',
 'anscombe',
 'attention',
 'brain_networks',
 'car_crashes',
 'diamonds',
 'dots',
 'dowjones',
 'exercise',
 'flights',
 'fmri',
 'geyser',
 'glue',
 'healthexp',
 'iris',
 'mpg',
 'penguins',
 'planets',
 'seaice',
 'taxis',
 'tips',
 'titanic']
```

In [91]: `# Load dataset`

```
sns.load_dataset('planets')
```

Out[91]:

| | method | number | orbital_period | mass | distance | year |
|------|-----------------|--------|----------------|-------|----------|------|
| 0 | Radial Velocity | 1 | 269.300000 | 7.10 | 77.40 | 2006 |
| 1 | Radial Velocity | 1 | 874.774000 | 2.21 | 56.95 | 2008 |
| 2 | Radial Velocity | 1 | 763.000000 | 2.60 | 19.84 | 2011 |
| 3 | Radial Velocity | 1 | 326.030000 | 19.40 | 110.62 | 2007 |
| 4 | Radial Velocity | 1 | 516.220000 | 10.50 | 119.47 | 2009 |
| ... | ... | ... | ... | ... | ... | ... |
| 1030 | Transit | 1 | 3.941507 | NaN | 172.00 | 2006 |
| 1031 | Transit | 1 | 2.615864 | NaN | 148.00 | 2007 |
| 1032 | Transit | 1 | 3.191524 | NaN | 174.00 | 2007 |
| 1033 | Transit | 1 | 4.125083 | NaN | 293.00 | 2008 |
| 1034 | Transit | 1 | 4.187757 | NaN | 260.00 | 2008 |

1035 rows × 6 columns

```
In [92]: sns.load_dataset('diamonds')
```

Out[92]:

| | carat | cut | color | clarity | depth | table | price | x | y | z |
|-------|-------|-----------|-------|---------|-------|-------|-------|------|------|------|
| 0 | 0.23 | Ideal | E | SI2 | 61.5 | 55.0 | 326 | 3.95 | 3.98 | 2.43 |
| 1 | 0.21 | Premium | E | SI1 | 59.8 | 61.0 | 326 | 3.89 | 3.84 | 2.31 |
| 2 | 0.23 | Good | E | VS1 | 56.9 | 65.0 | 327 | 4.05 | 4.07 | 2.31 |
| 3 | 0.29 | Premium | I | VS2 | 62.4 | 58.0 | 334 | 4.20 | 4.23 | 2.63 |
| 4 | 0.31 | Good | J | SI2 | 63.3 | 58.0 | 335 | 4.34 | 4.35 | 2.75 |
| ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... |
| 53935 | 0.72 | Ideal | D | SI1 | 60.8 | 57.0 | 2757 | 5.75 | 5.76 | 3.50 |
| 53936 | 0.72 | Good | D | SI1 | 63.1 | 55.0 | 2757 | 5.69 | 5.75 | 3.61 |
| 53937 | 0.70 | Very Good | D | SI1 | 62.8 | 60.0 | 2757 | 5.66 | 5.68 | 3.56 |
| 53938 | 0.86 | Premium | H | SI2 | 61.0 | 58.0 | 2757 | 6.15 | 6.12 | 3.74 |
| 53939 | 0.75 | Ideal | D | SI2 | 62.2 | 55.0 | 2757 | 5.83 | 5.87 | 3.64 |

53940 rows × 10 columns

```
In [93]: sns.load_dataset('healthexp')
```

Out[93]:

| | Year | Country | Spending_USD | Life_Expectancy |
|-----|------|---------------|--------------|-----------------|
| 0 | 1970 | Germany | 252.311 | 70.6 |
| 1 | 1970 | France | 192.143 | 72.2 |
| 2 | 1970 | Great Britain | 123.993 | 71.9 |
| 3 | 1970 | Japan | 150.437 | 72.0 |
| 4 | 1970 | USA | 326.961 | 70.9 |
| ... | ... | ... | ... | ... |
| 269 | 2020 | Germany | 6938.983 | 81.1 |
| 270 | 2020 | France | 5468.418 | 82.3 |
| 271 | 2020 | Great Britain | 5018.700 | 80.4 |
| 272 | 2020 | Japan | 4665.641 | 84.7 |
| 273 | 2020 | USA | 11859.179 | 77.0 |

274 rows × 4 columns

```
In [ ]:
```

