

```
In [1]: import seaborn as sns
```

```
In [2]: df = sns.load_dataset("tips")
df.head()
```

Out[2]:

	total_bill	tip	sex	smoker	day	time	size
0	16.99	1.01	Female	No	Sun	Dinner	2
1	10.34	1.66	Male	No	Sun	Dinner	3
2	21.01	3.50	Male	No	Sun	Dinner	3
3	23.68	3.31	Male	No	Sun	Dinner	2
4	24.59	3.61	Female	No	Sun	Dinner	4

## Regression Plot

- It creates a regression line between two parameters and helps to visualize their linear relationship
- lmplot

```
In [3]: print(dir(sns))
```

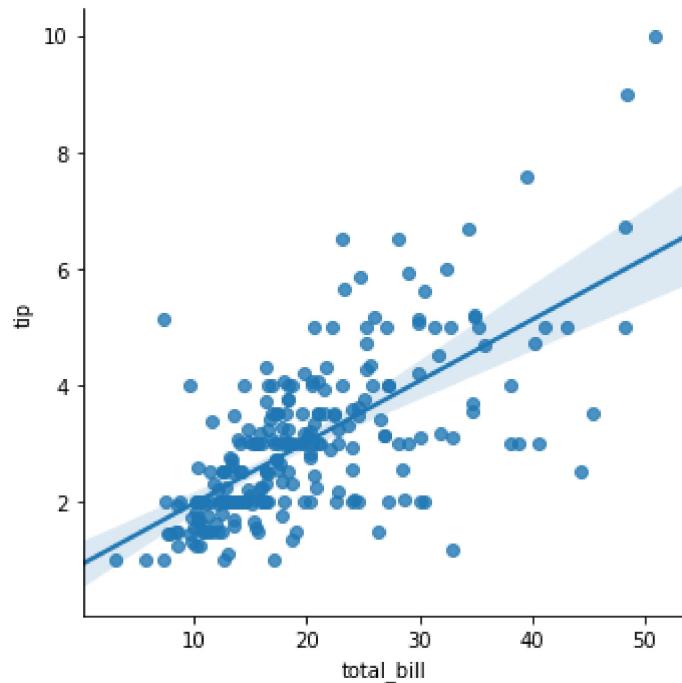
```
['FacetGrid', 'JointGrid', 'PairGrid', '__builtins__', '__cached__', '__doc__', '__file__', '__loader__', '__name__', '__package__', '__path__', '__spec__', '__version__', '_orig_rc_params', 'algorithms', 'axes_style', 'axisgrid', 'barplot', 'blend_palette', 'boxenplot', 'boxplot', 'categorical', 'catplot', 'choose_colorbrewer_palette', 'choose_cubehelix_palette', 'choose_dark_palette', 'choose_diverging_palette', 'choose_light_palette', 'clustermap', 'cm', 'color_palette', 'colors', 'countplot', 'crayon_palette', 'crayons', 'cubehelix_palette', 'dark_palette', 'desaturate', 'despine', 'distplot', 'distributions', 'diverging_palette', 'dogplot', 'external', 'factorplot', 'get_dataset_names', 'heatmap', 'hls_palette', 'husl_palette', 'jointplot', 'kdeplot', 'light_palette', 'lineplot', 'lmplot', 'load_dataset', 'lvplot', 'matrix', 'miscplot', 'mpl', 'mpl_palette', 'pairplot', 'palettes', 'palplot', 'plotting_context', 'pointplot', 'rcmod', 'regplot', 'regression', 'relational', 'relplot', 'reset_defaults', 'reset_orig', 'residplot', 'rugplot', 'saturate', 'scatterplot', 'set', 'set_color_codes', 'set_context', 'set_hls_values', 'set_palette', 'set_style', 'stripplot', 'swarmplot', 'timeseries', 'tsplot', 'utils', 'violinplot', 'widgets', 'xkcd_palette', 'xkcd_rgb']
```

```
In [4]: df.columns
```

```
Out[4]: Index(['total_bill', 'tip', 'sex', 'smoker', 'day', 'time', 'size'], dtype='object')
```

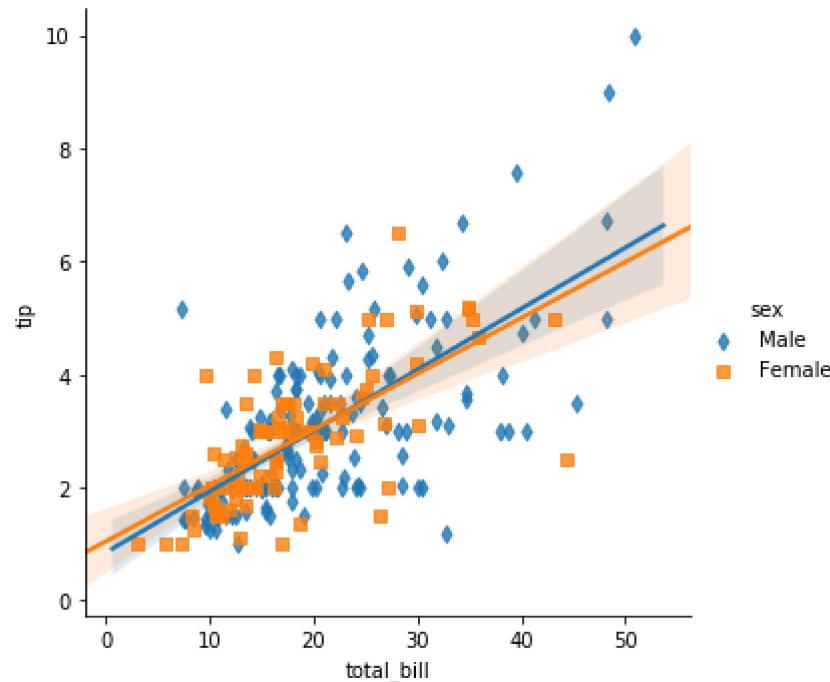
```
In [5]: sns.lmplot(x="total_bill",y="tip",data=df)
```

```
Out[5]: <seaborn.axisgrid.FacetGrid at 0x14a7d40aa90>
```



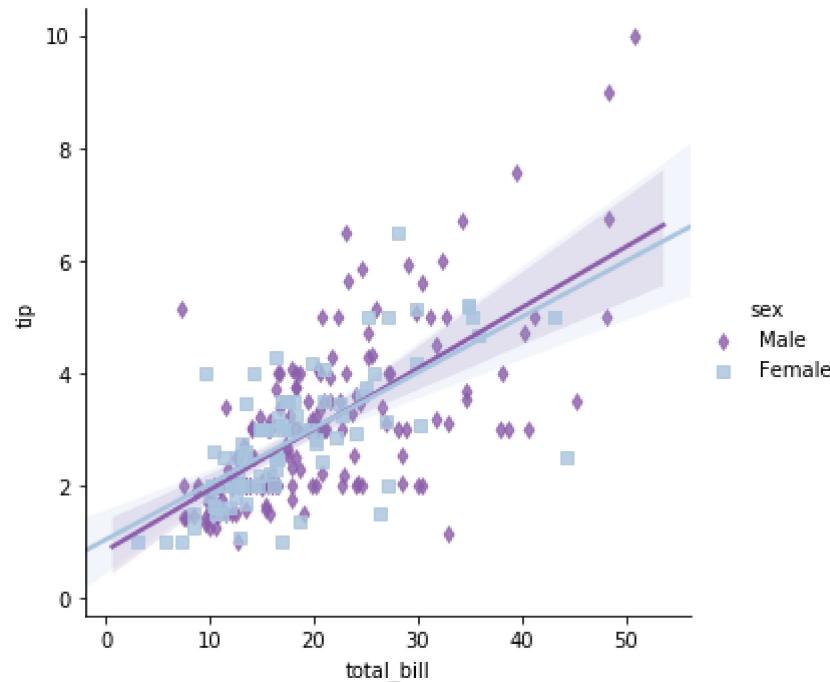
```
In [7]: sns.lmplot(x="total_bill",y="tip",data=df,hue="sex",markers=["d","s"])
```

```
Out[7]: <seaborn.axisgrid.FacetGrid at 0x14a7e7ed940>
```



```
In [13]: sns.lmplot(x="total_bill",y="tip",data=df,hue="sex",markers=["d","s"],palette="BuPu_r",scatter_kws={"s":30})
```

```
Out[13]: <seaborn.axisgrid.FacetGrid at 0x14a7ece83c8>
```



In [14]: `help(sns.lmplot)`

```
e, x_estimator=None, x_bins=None, x_ci='ci', scatter=True, fit_reg=True, ci=95, n_boot=1000, units=None, order=1, logistic=False, lowess=False, robust=False, logx=False, x_partial=None, y_partial=None, truncate=False, x_jitter=None, y_jitter=None, scatter_kws=None, line_kws=None, size=None)
    Plot data and regression model fits across a FacetGrid.
```

This function combines :func:`regplot` and :class:`FacetGrid`. It is intended as a convenient interface to fit regression models across conditional subsets of a dataset.

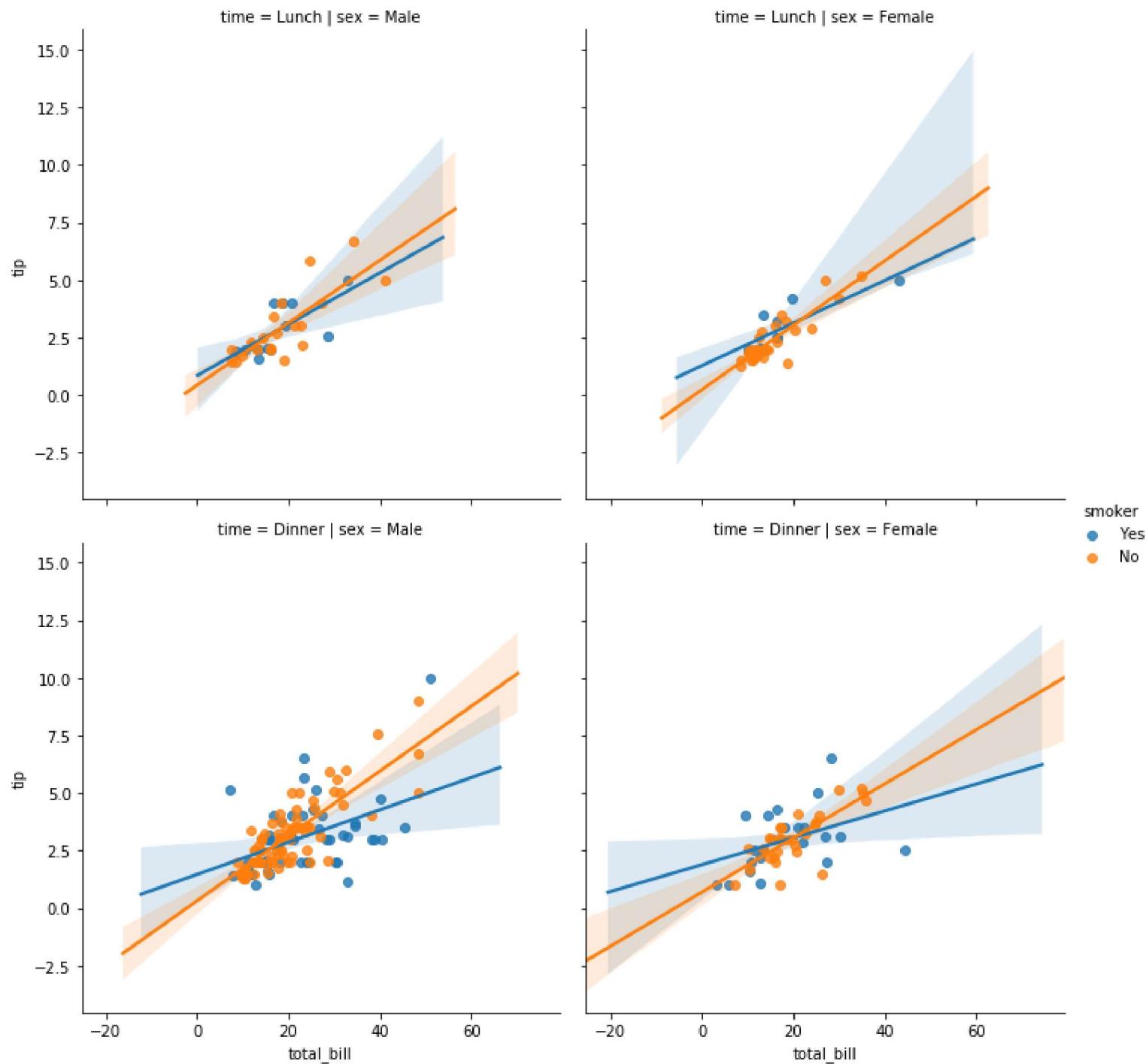
When thinking about how to assign variables to different facets, a general rule is that it makes sense to use ``hue`` for the most important comparison, followed by ``col`` and ``row``. However, always think about your particular dataset and the goals of the visualization you are creating.

There are a number of mutually exclusive options for estimating the regression model. See the :ref:`tutorial <regression\_tutorial>` for more information.

The parameters to this function span most of the options in

```
In [15]: # Displaying Multiple plots  
sns.lmplot(x="total_bill",y="tip",data=df,col="sex",row="time",hue="smoker")
```

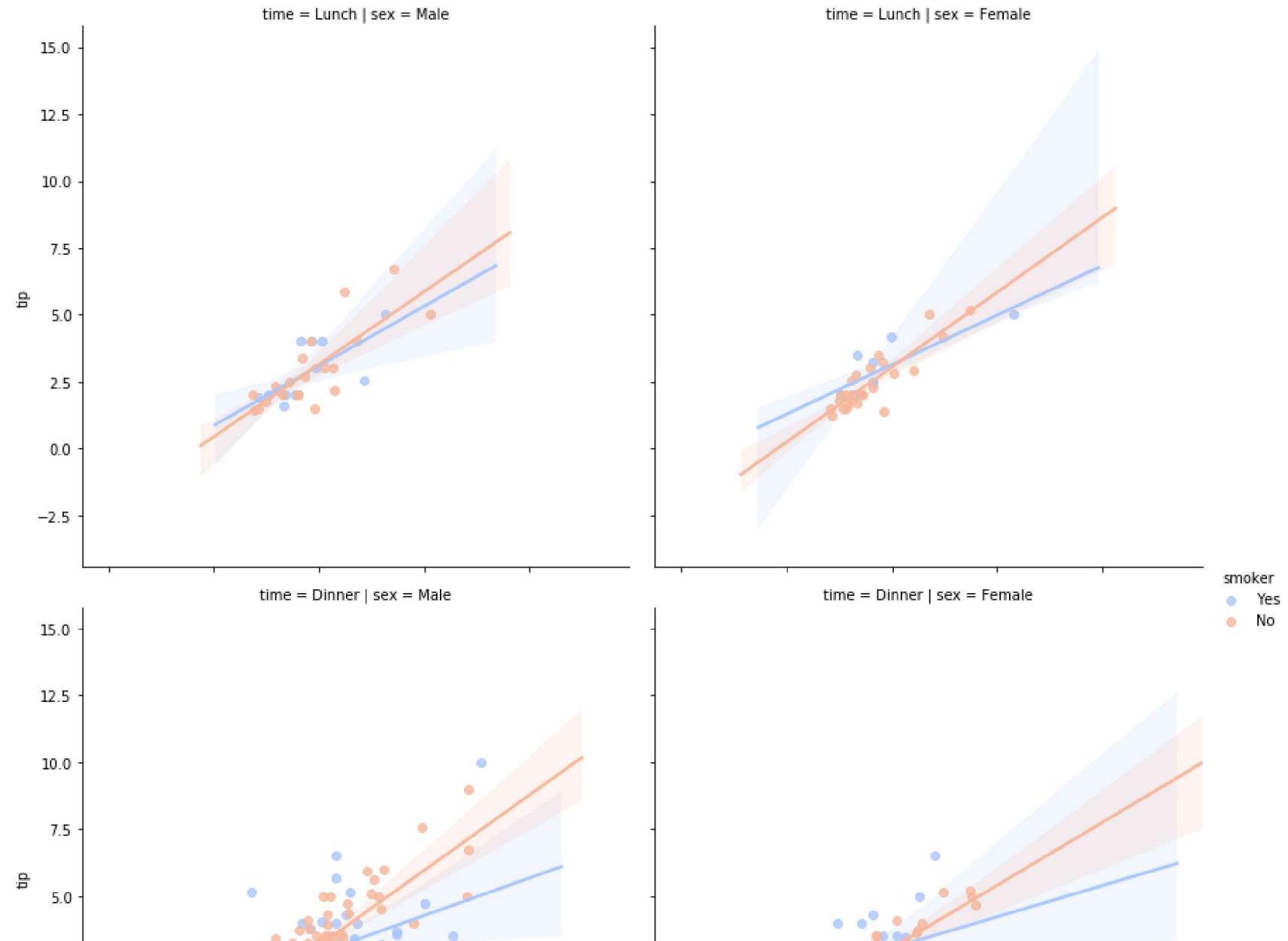
```
Out[15]: <seaborn.axisgrid.FacetGrid at 0x14a7edb0550>
```

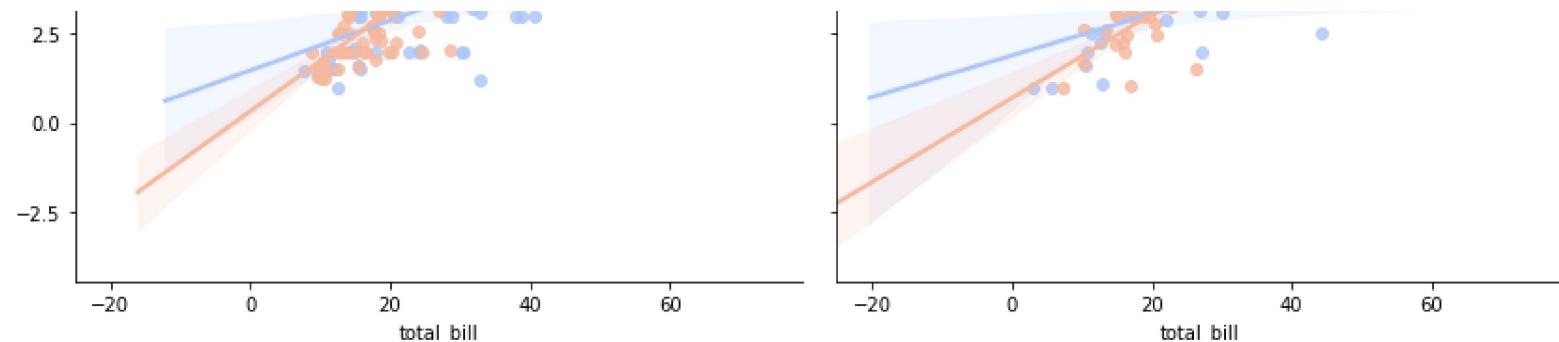


In [16]: # Displaying Multiple plots

```
sns.lmplot(x="total_bill",y="tip",data=df,col="sex",row="time",hue="smoker",height=6,palette="coolwarm")
```

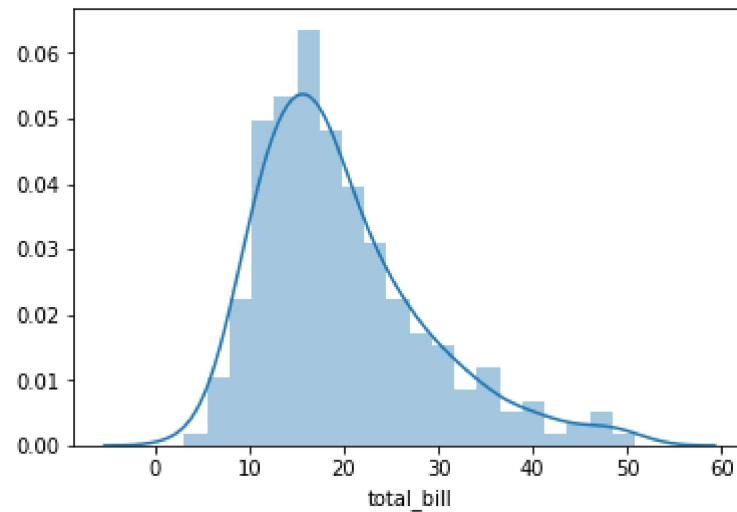
Out[16]: <seaborn.axisgrid.FacetGrid at 0x14a7f181978>





```
In [24]: # bin represent how many falls in data  
sns.distplot(df["total_bill"],bins=20)
```

```
Out[24]: <matplotlib.axes._subplots.AxesSubplot at 0x14a7fa3f1d0>
```



```
In [ ]: # Matrix plot and heatmap
```

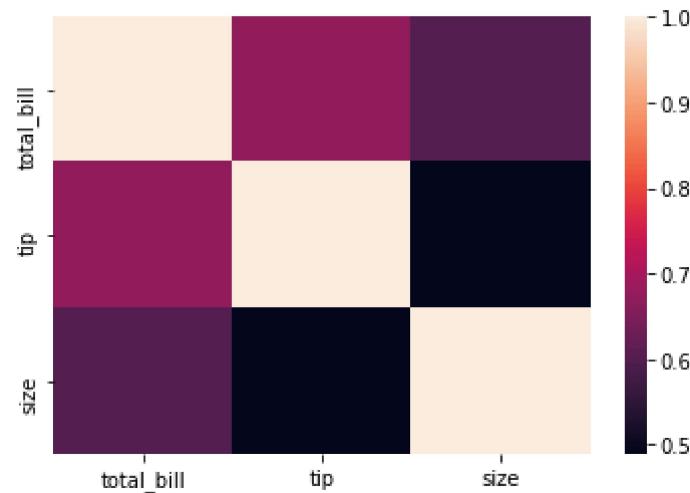
```
In [25]: tc = df.corr()  
tc
```

Out[25]:

	total_bill	tip	size
total_bill	1.000000	0.675734	0.598315
tip	0.675734	1.000000	0.489299
size	0.598315	0.489299	1.000000

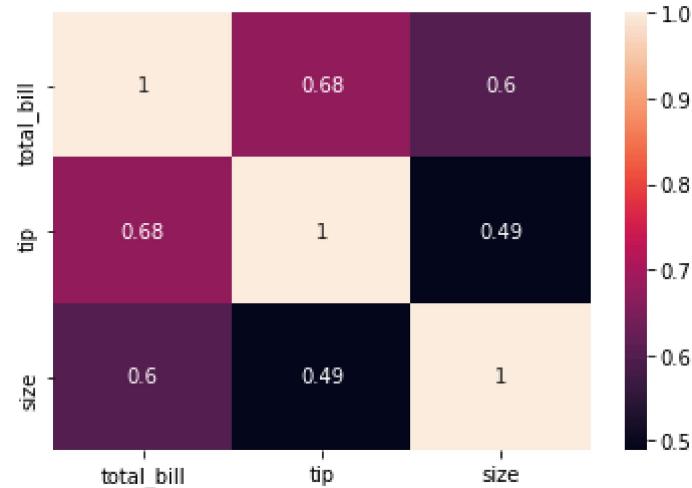
```
In [26]: sns.heatmap(tc)
```

Out[26]: <matplotlib.axes.\_subplots.AxesSubplot at 0x14a7fb28208>



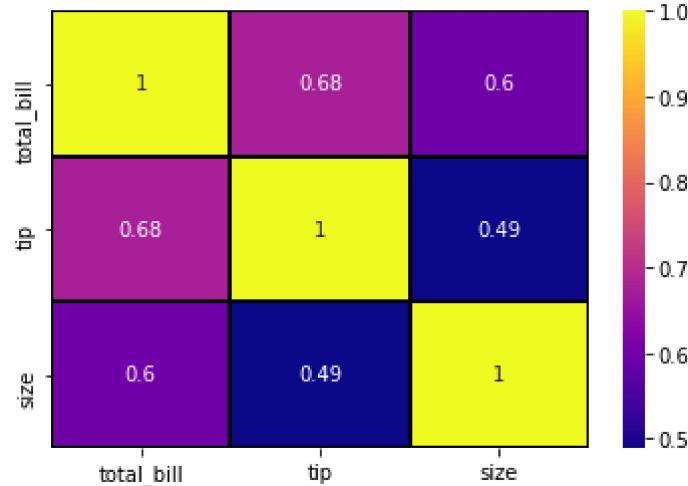
```
In [27]: sns.heatmap(tc, annot=True)
```

```
Out[27]: <matplotlib.axes._subplots.AxesSubplot at 0x14a7fbc1780>
```



```
In [28]: sns.heatmap(tc, annot=True, cmap="plasma", linecolor="black", linewidth=1)
```

```
Out[28]: <matplotlib.axes._subplots.AxesSubplot at 0x14a7ff8d208>
```



```
In [30]: sns.get_dataset_names()
```

C:\Users\Alekhya\Anaconda3\lib\site-packages\seaborn\utils.py:376: UserWarning: No parser was explicitly specified, so I'm using the best available HTML parser for this system ("lxml"). This usually isn't a problem, but if you run this code on another system, or in a different virtual environment, it may use a different parser and behave differently.

The code that caused this warning is on line 376 of the file C:\Users\Alekhya\Anaconda3\lib\site-packages\seaborn\utils.py. To get rid of this warning, pass the additional argument 'features="lxml"' to the BeautifulSoup constructor.

```
gh_list = BeautifulSoup(http)
```

```
Out[30]: ['anagrams',
 'anscombe',
 'attention',
 'brain_networks',
 'car_crashes',
 'diamonds',
 'dots',
 'exercise',
 'flights',
 'fmri',
 'gammas',
 'geyser',
 'iris',
 'mpg',
 'penguins',
 'planets',
 'tips',
 'titanic']
```

```
In [29]: s = sns.load_dataset("iris")
s.head()
```

Out[29]:

	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

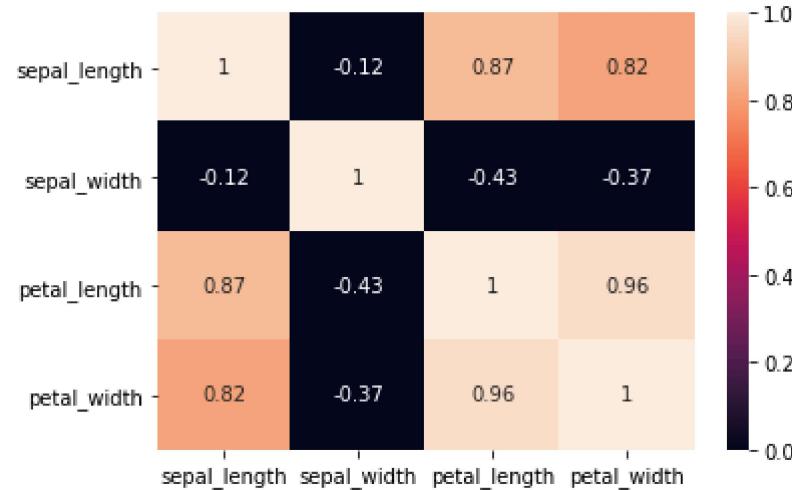
```
In [31]: s1 = s.corr()
s1
```

Out[31]:

	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

```
In [58]: sns.heatmap(s1, annot=True, vmin=0, vmax=1)
```

```
Out[58]: <matplotlib.axes._subplots.AxesSubplot at 0x14a03c4c2b0>
```



```
In [37]: help(sns.heatmap)
```

```
-----  
data : rectangular dataset  
    2D dataset that can be coerced into an ndarray. If a Pandas DataFrame  
    is provided, the index/column information will be used to label the  
    columns and rows.  
vmin, vmax : floats, optional  
    Values to anchor the colormap, otherwise they are inferred from the  
    data and other keyword arguments.  
cmap : matplotlib colormap name or object, or list of colors, optional  
    The mapping from data values to color space. If not provided, the  
    default will depend on whether ``center`` is set.  
center : float, optional  
    The value at which to center the colormap when plotting divergent data.  
    Using this parameter will change the default ``cmap`` if none is  
    specified.  
robust : bool, optional  
    If True and ``vmin`` or ``vmax`` are absent, the colormap range is  
    computed with robust quantiles instead of the extreme values.  
annot : bool or rectangular dataset, optional  
    If True, write the data value in each cell. If an array-like with the
```

```
In [38]: flights = sns.load_dataset("flights")  
flights.head()
```

Out[38]:

	year	month	passengers
0	1949	January	112
1	1949	February	118
2	1949	March	132
3	1949	April	129
4	1949	May	121

```
In [44]: f = flights.pivot("year", "month", "passengers")
f.head()
```

Out[44]:

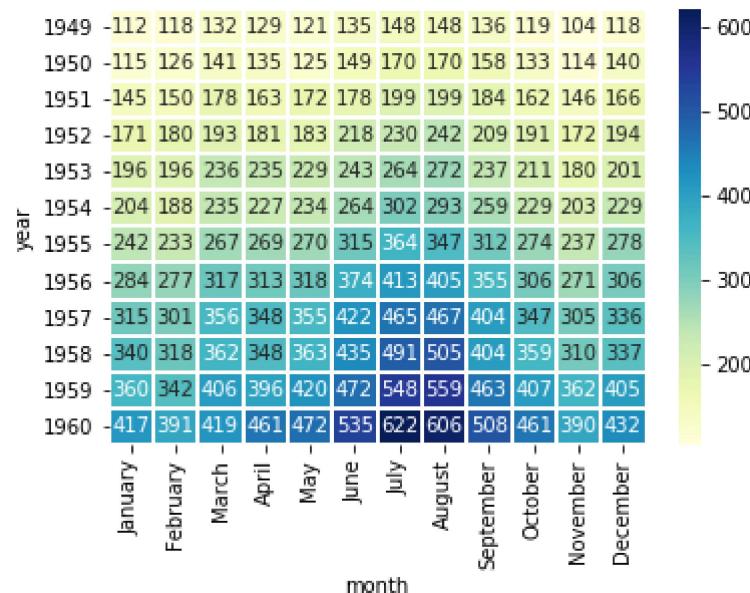
month	January	February	March	April	May	June	July	August	September	October	November	December
year												
1949	112	118	132	129	121	135	148	148	136	119	104	118
1950	115	126	141	135	125	149	170	170	158	133	114	140
1951	145	150	178	163	172	178	199	199	184	162	146	166
1952	171	180	193	181	183	218	230	242	209	191	172	194
1953	196	196	236	235	229	243	264	272	237	211	180	201

```
In [45]: f.shape
```

Out[45]: (12, 12)

```
In [48]: sns.heatmap(f, annot=True, cmap="YlGnBu", linewidths=1, fmt="d")
```

```
Out[48]: <matplotlib.axes._subplots.AxesSubplot at 0x14a02b3db38>
```



```
In [49]: f1 = f.corr()  
f1
```

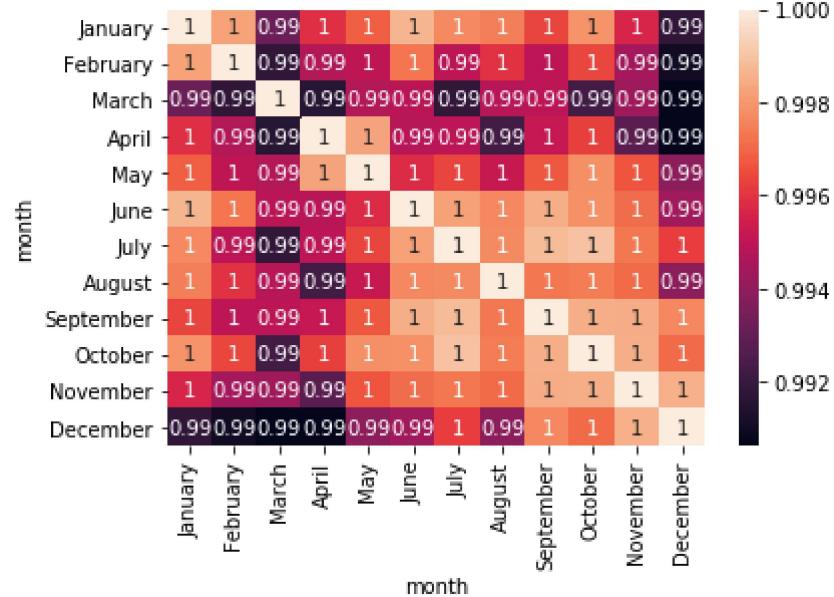
Out[49]:

month	January	February	March	April	May	June	July	August	September	October	November	December
month												
January	1.000000	0.998247	0.993215	0.995918	0.996823	0.998700	0.997717	0.997479	0.996372	0.997920	0.995648	0.991821
February	0.998247	1.000000	0.991813	0.994752	0.995039	0.997296	0.994991	0.995954	0.995011	0.996401	0.994475	0.991053
March	0.993215	0.991813	1.000000	0.991589	0.994742	0.994923	0.991882	0.994852	0.994969	0.992312	0.994334	0.990635
April	0.995918	0.994752	0.991589	1.000000	0.998290	0.994910	0.994981	0.992308	0.995188	0.996277	0.992867	0.990684
May	0.996823	0.995039	0.994742	0.998290	1.000000	0.995804	0.996357	0.995219	0.996724	0.997837	0.996645	0.993936
June	0.998700	0.997296	0.994923	0.994910	0.995804	1.000000	0.998211	0.997674	0.998527	0.997856	0.997113	0.994374
July	0.997717	0.994991	0.991882	0.994981	0.996357	0.998211	1.000000	0.997756	0.998808	0.998987	0.997343	0.996202
August	0.997479	0.995954	0.994852	0.992308	0.995219	0.997674	0.997756	1.000000	0.997334	0.997429	0.997102	0.993931
September	0.996372	0.995011	0.994969	0.995188	0.996724	0.998527	0.998808	0.997334	1.000000	0.998562	0.998478	0.997680
October	0.997920	0.996401	0.992312	0.996277	0.997837	0.997856	0.998987	0.997429	0.998562	1.000000	0.998563	0.997108
November	0.995648	0.994475	0.994334	0.992867	0.996645	0.997113	0.997343	0.997102	0.998478	0.998563	1.000000	0.998601
December	0.991821	0.991053	0.990635	0.990684	0.993936	0.994374	0.996202	0.993931	0.997680	0.997108	0.998601	1.000000



```
In [50]: sns.heatmap(f1,annot=True)
```

```
Out[50]: <matplotlib.axes._subplots.AxesSubplot at 0x14a013dc668>
```



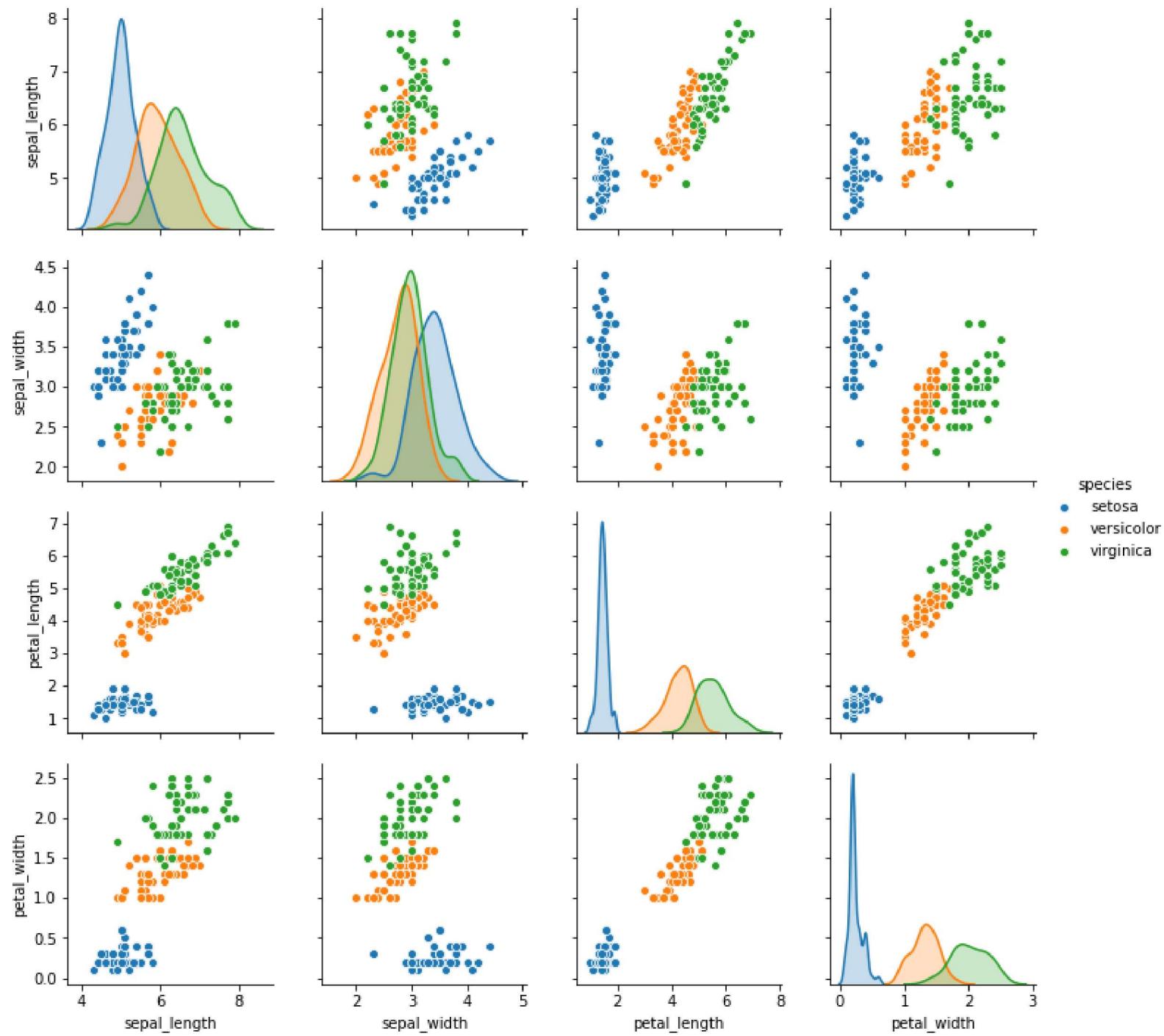
In [53]: `s.head()`

Out[53]:

	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

```
In [52]: #pairplot  
sns.pairplot(s,hue="species")
```

```
Out[52]: <seaborn.axisgrid.PairGrid at 0x14a01491cf8>
```



```
In [54]: help(sns.heatmap)
```

```
o', yticklabels='auto', mask=None, ax=None, **kwargs)
Plot rectangular data as a color-encoded matrix.
```

This is an Axes-level function and will draw the heatmap into the currently-active Axes if none is provided to the ``ax`` argument. Part of this Axes space will be taken and used to plot a colormap, unless ``cbar`` is False or a separate Axes is provided to ``cbar\_ax``.

#### Parameters

-----

`data` : rectangular dataset

2D dataset that can be coerced into an ndarray. If a Pandas DataFrame is provided, the index/column information will be used to label the columns and rows.

`vmin`, `vmax` : floats, optional

Values to anchor the colormap, otherwise they are inferred from the data and other keyword arguments.

`cmap` : matplotlib colormap name or object, or list of colors, optional

The mapping from data values to color space. If not provided, the default will depend on whether ``center`` is set.

```
In [ ]:
```

