

In [1]:

```
import seaborn as sn
```

In [3]:

```
iris = sn.load_dataset("iris")
```

In [4]:

```
iris
```

Out[4]:

|     | 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    |
| ... | ...          | ...         | ...          | ...         | ...       |
| 145 | 6.7          | 3.0         | 5.2          | 2.3         | virginica |
| 146 | 6.3          | 2.5         | 5.0          | 1.9         | virginica |
| 147 | 6.5          | 3.0         | 5.2          | 2.0         | virginica |
| 148 | 6.2          | 3.4         | 5.4          | 2.3         | virginica |
| 149 | 5.9          | 3.0         | 5.1          | 1.8         | virginica |

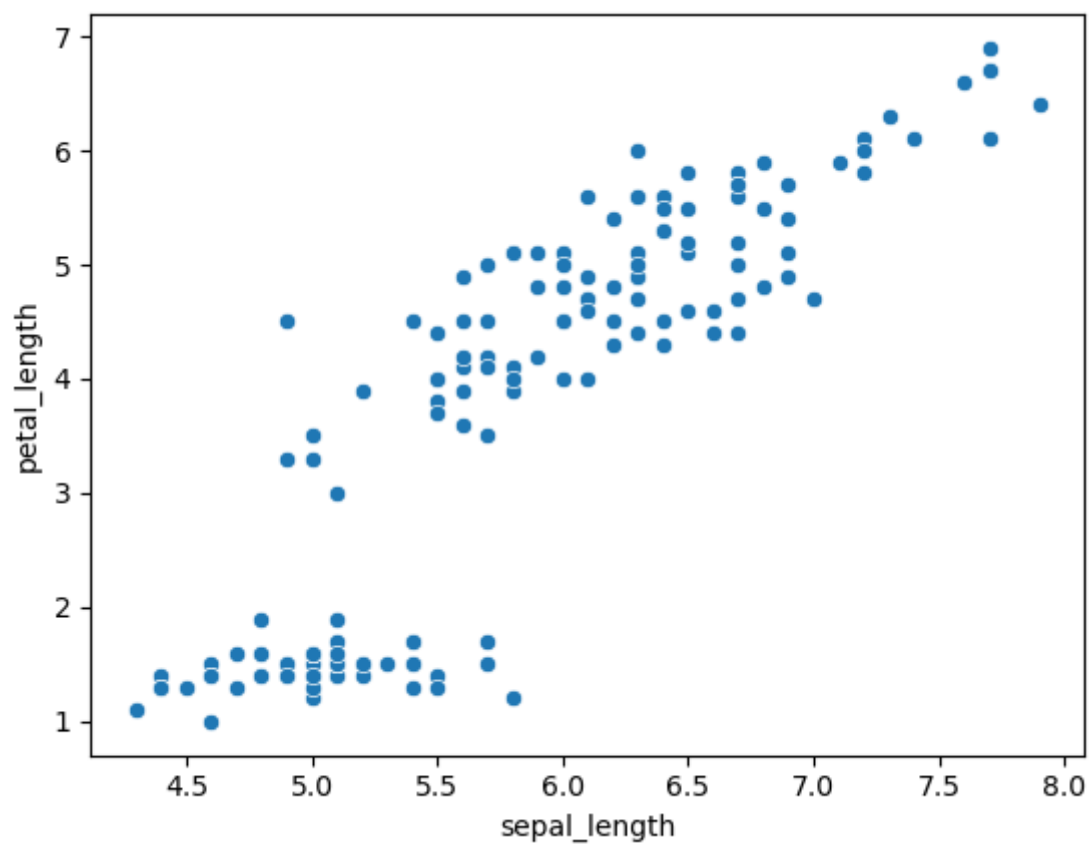
150 rows × 5 columns

In [7]:

```
sn.scatterplot(x= iris.sepal_length,y = iris.petal_length)
```

Out[7]:

<Axes: xlabel='sepal\_length', ylabel='petal\_length'>

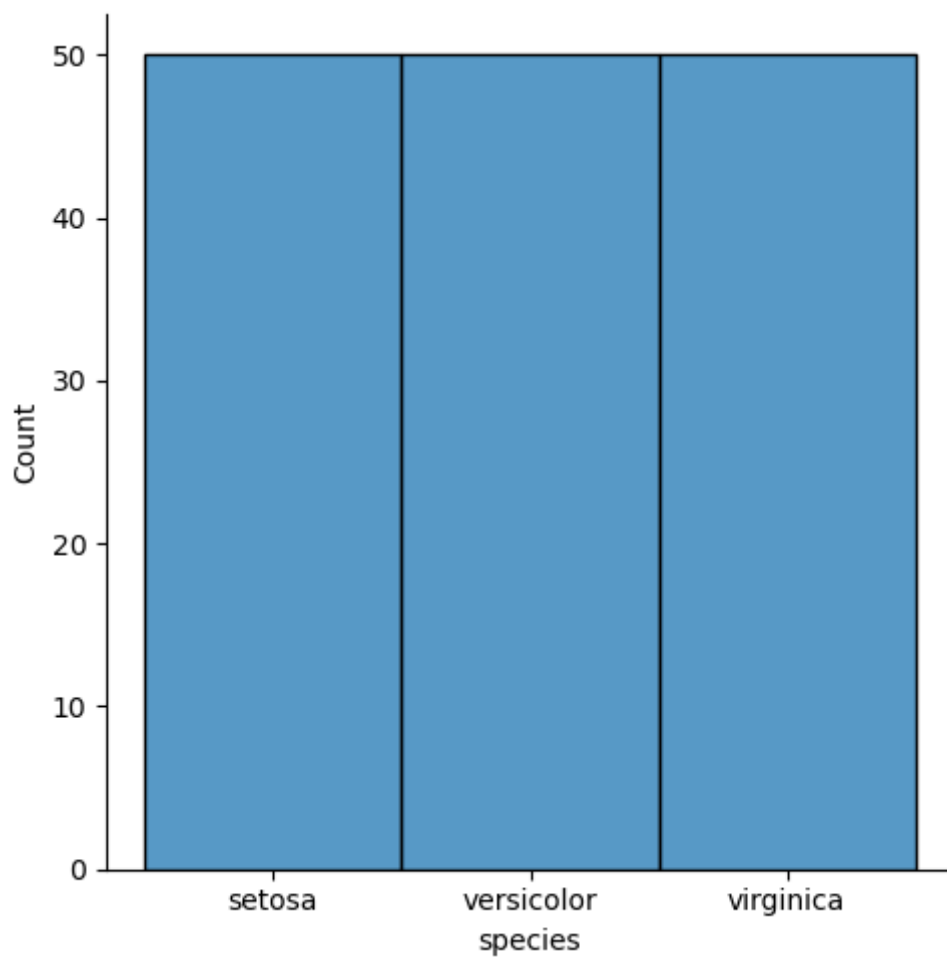


In [9]:

```
sn.displot(iris["species"])
```

Out[9]:

<seaborn.axisgrid.FacetGrid at 0x1f93d5d4460>

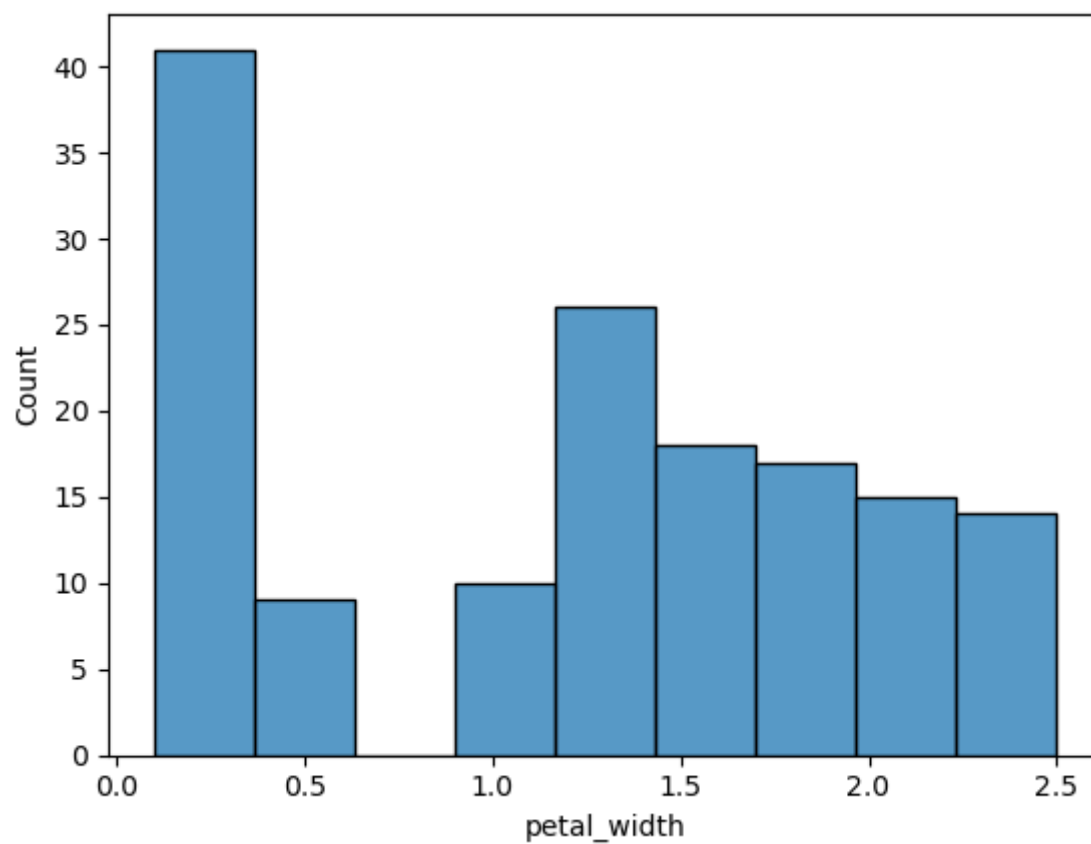


In [11]:

```
sn.histplot(iris.petal_width)
```

Out[11]:

<Axes: xlabel='petal\_width', ylabel='Count'>



In [12]:

```
tips = sn.load_dataset("tips")
```

In [13]:

tips

Out[13]:

|     | 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    |
| ... | ...        | ...  | ...    | ...    | ...  | ...    | ...  |
| 239 | 29.03      | 5.92 | Male   | No     | Sat  | Dinner | 3    |
| 240 | 27.18      | 2.00 | Female | Yes    | Sat  | Dinner | 2    |
| 241 | 22.67      | 2.00 | Male   | Yes    | Sat  | Dinner | 2    |
| 242 | 17.82      | 1.75 | Male   | No     | Sat  | Dinner | 2    |
| 243 | 18.78      | 3.00 | Female | No     | Thur | Dinner | 2    |

244 rows × 7 columns

In [17]:

tips["smoker"].value\_count()

-----  
 -

**AttributeError** Traceback (most recent call last)  
 t)

Cell In[17], line 1

----&gt; 1 tips["smoker"].value\_count()

File ~\anaconda3\lib\site-packages\pandas\core\generic.py:5902, in NDFrame

e.\_\_getattr\_\_(self, name)

```

5895 if (
5896     name not in self._internal_names_set
5897     and name not in self._metadata
5898     and name not in self._accessors
5899     and self._info_axis._can_hold_identifiers_and_holds_name(name)
5900 ):
5901     return self[name]
-> 5902 return object.__getattr__(self, name)

```

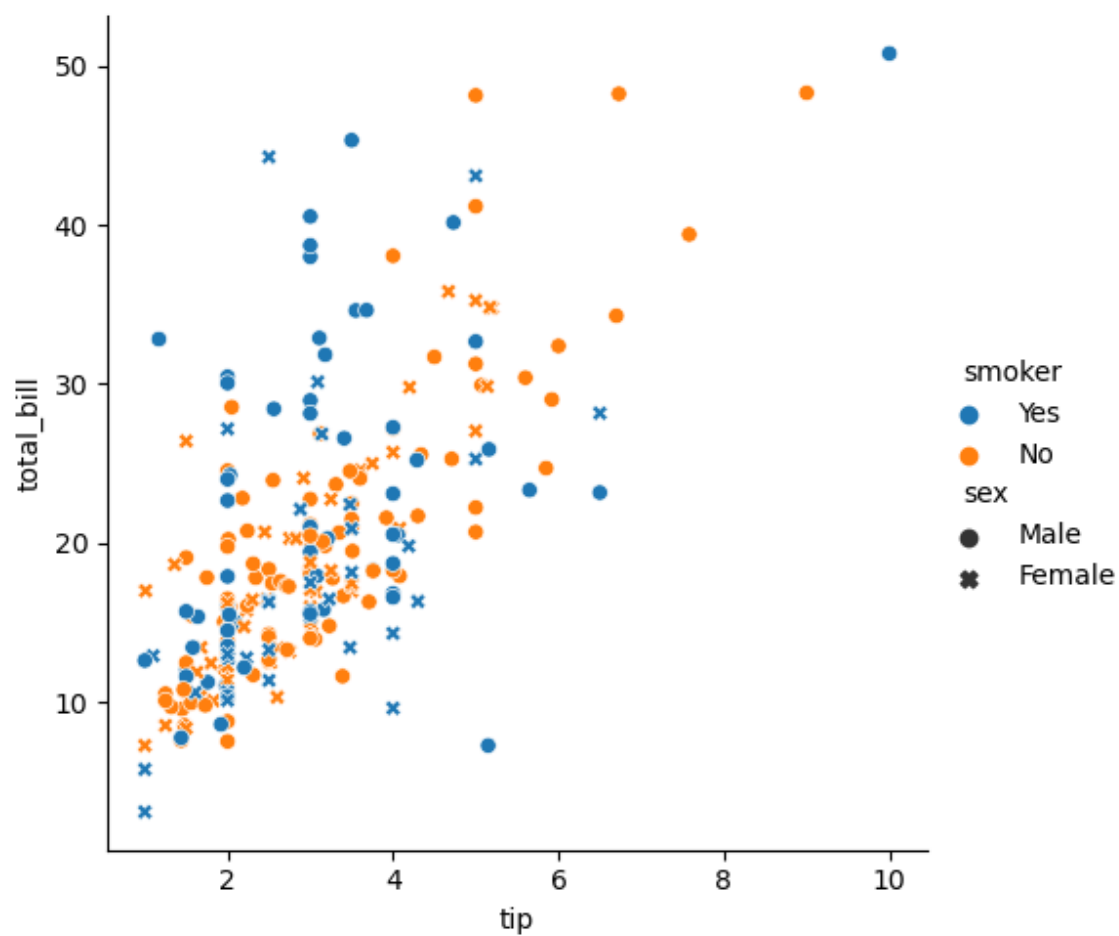
**AttributeError:** 'Series' object has no attribute 'value\_count'

In [22]:

```
sn.relplot(x =tips.tip,y = tips.total_bill,data= tips,style="sex",hue= "smoker" )
```

Out[22]:

<seaborn.axisgrid.FacetGrid at 0x1f944bfcca0>

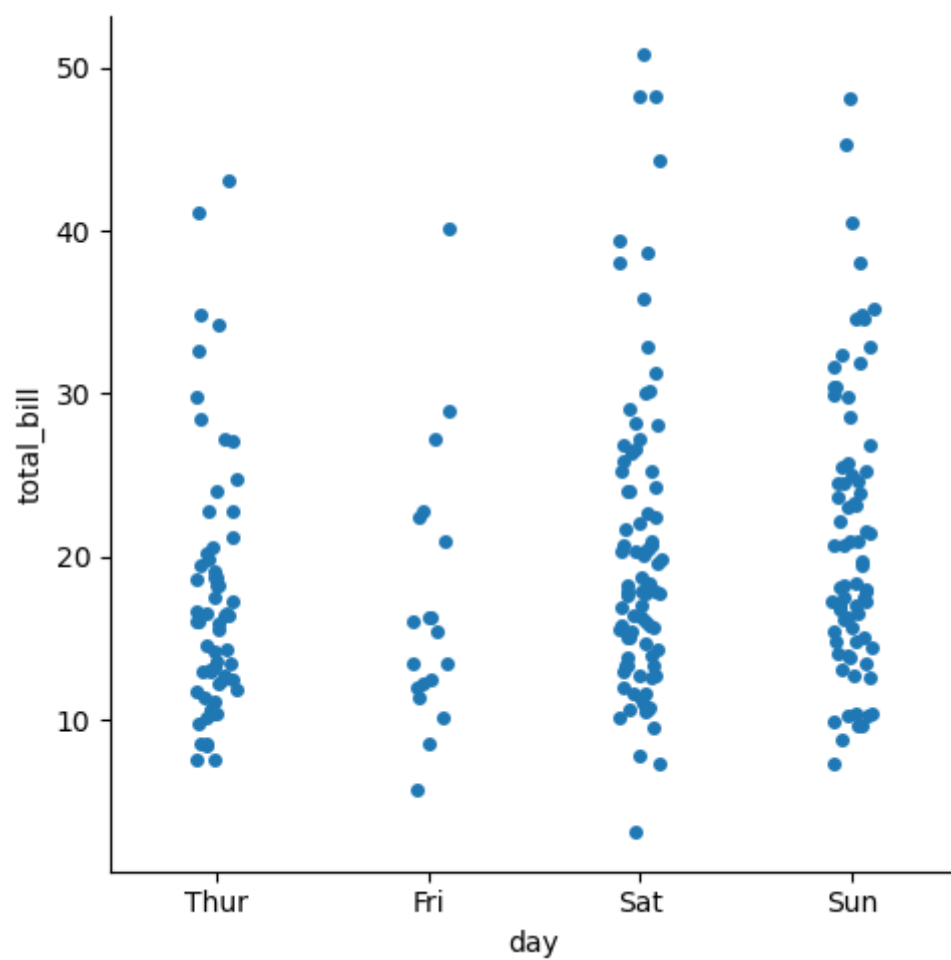


In [25]:

```
sn.catplot(x = tips.day,y = tips.total_bill,data= tips )
```

Out[25]:

<seaborn.axisgrid.FacetGrid at 0x1f944e575b0>

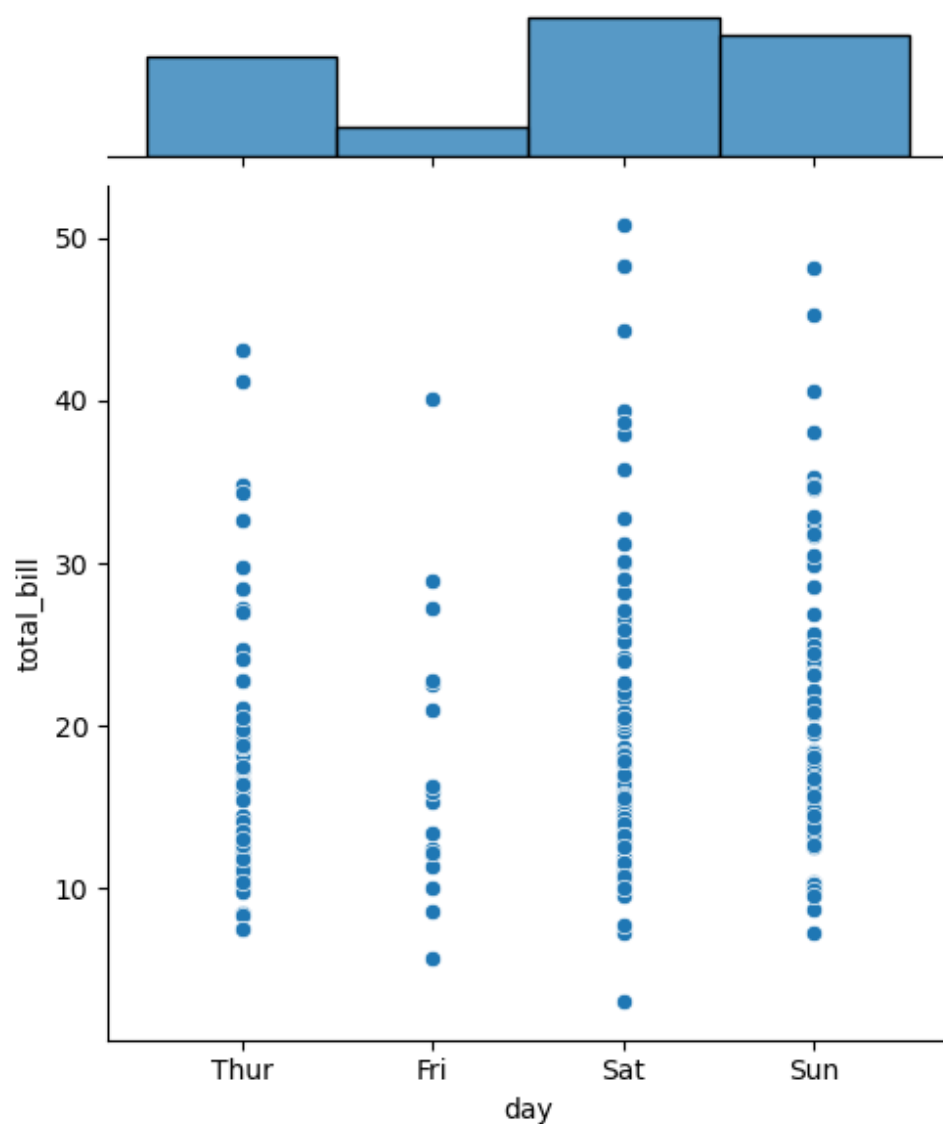


In [26]:

```
sn.jointplot(x = tips.day,y = tips.total_bill)
```

Out[26]:

<seaborn.axisgrid.JointGrid at 0x1f944e3fca0>



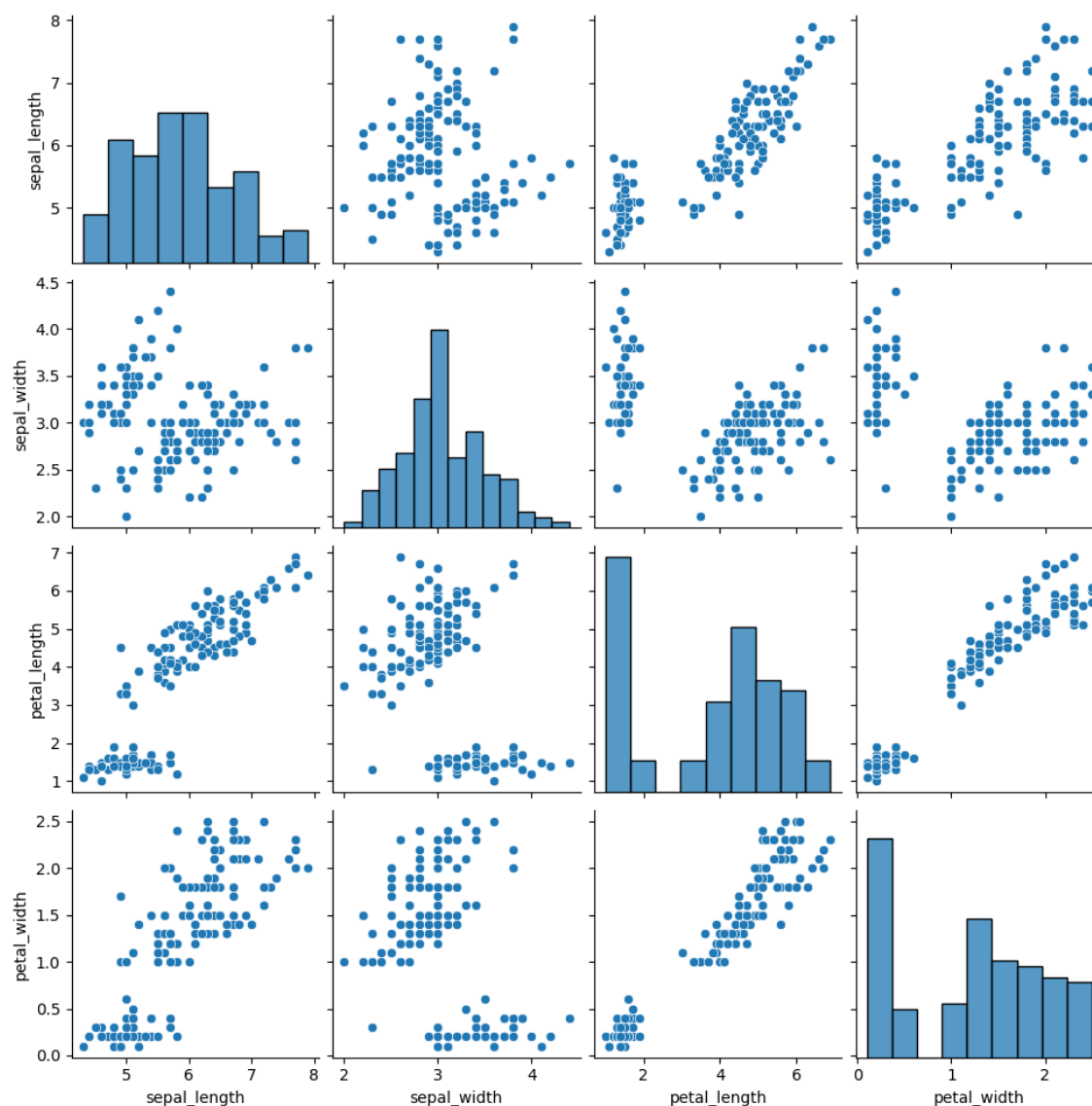


In [27]:

```
sn.pairplot(iris)
```

Out[27]:

&lt;seaborn.axisgrid.PairGrid at 0x1f94522fac0&gt;

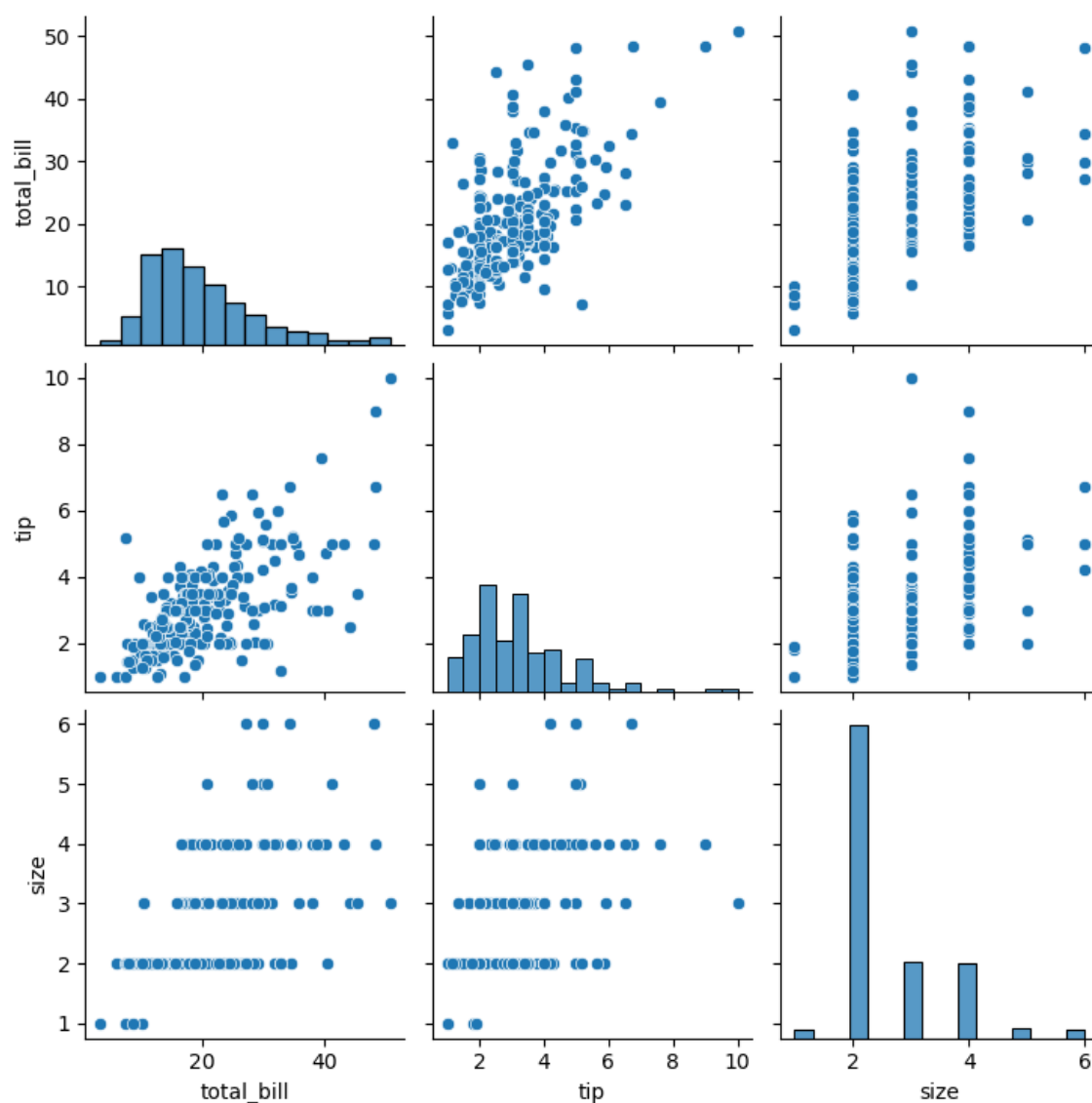


In [28]:

```
sn.pairplot(tips)
```

Out[28]:

<seaborn.axisgrid.PairGrid at 0x1f941a21090>

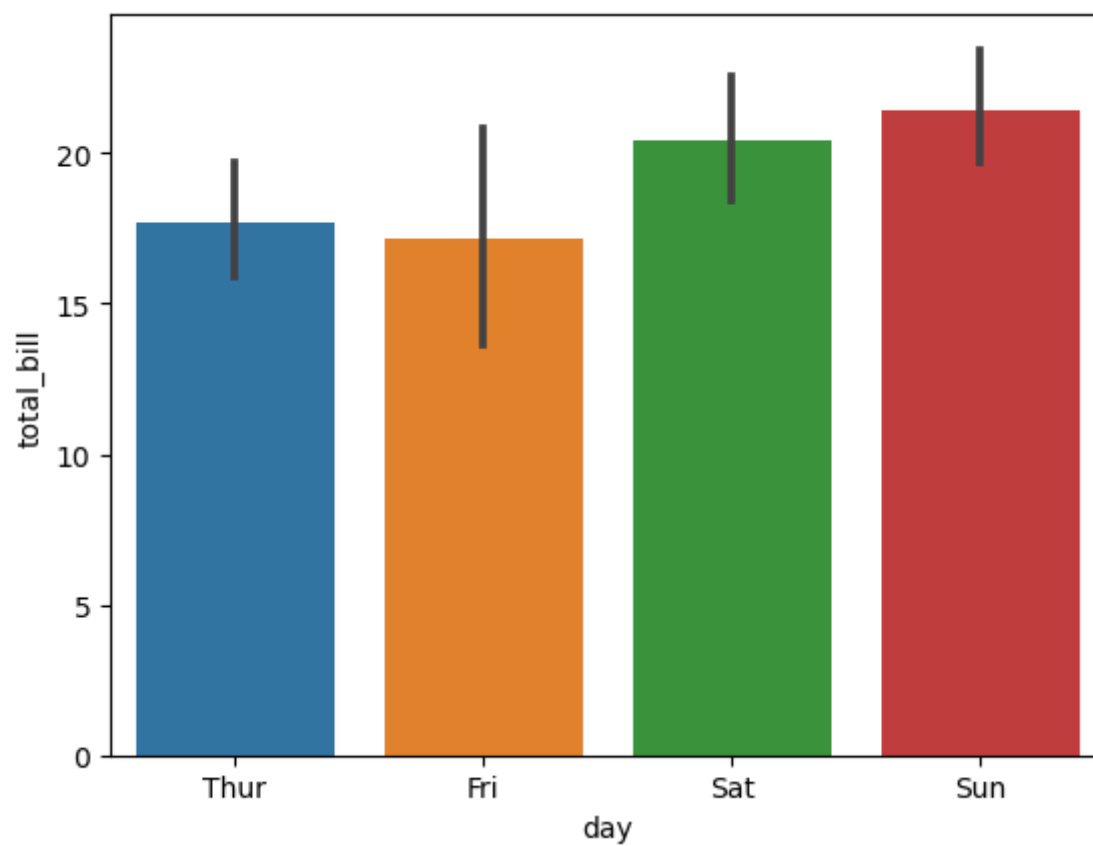


In [31]:

```
sn.barplot(x= tips.day,y = tips.total_bill)
```

Out[31]:

<Axes: xlabel='day', ylabel='total\_bill'>

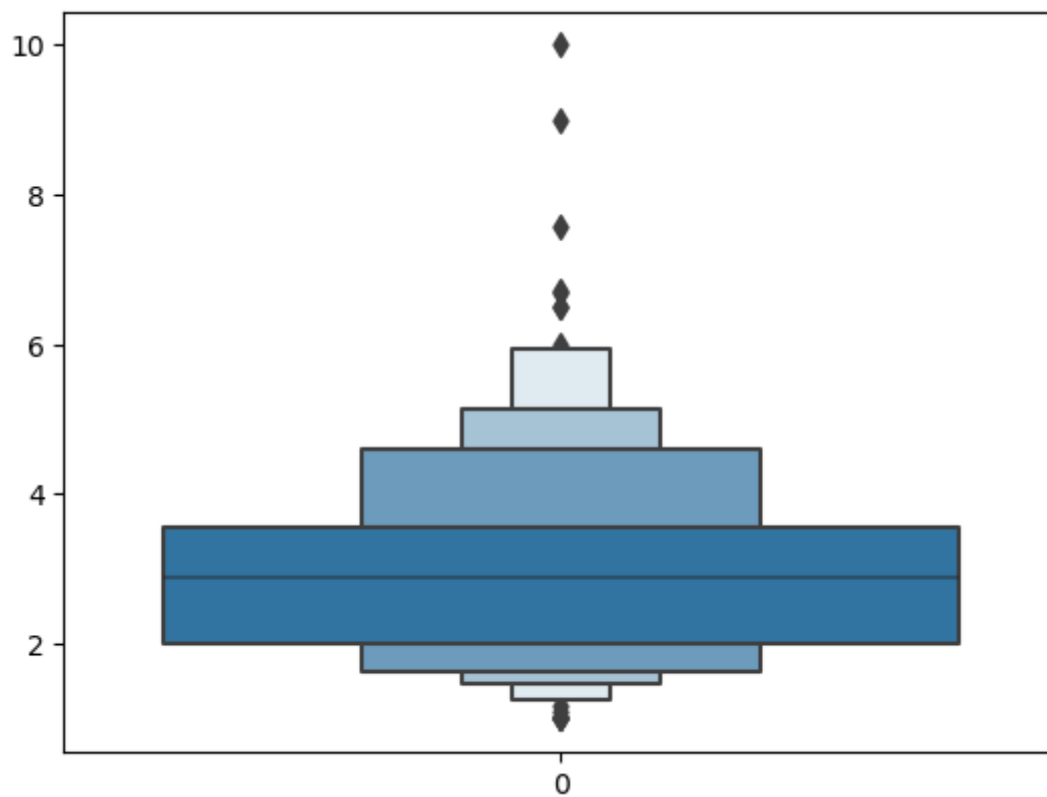


In [34]:

```
sn.boxenplot(tips.tip)
```

Out[34]:

<Axes: >

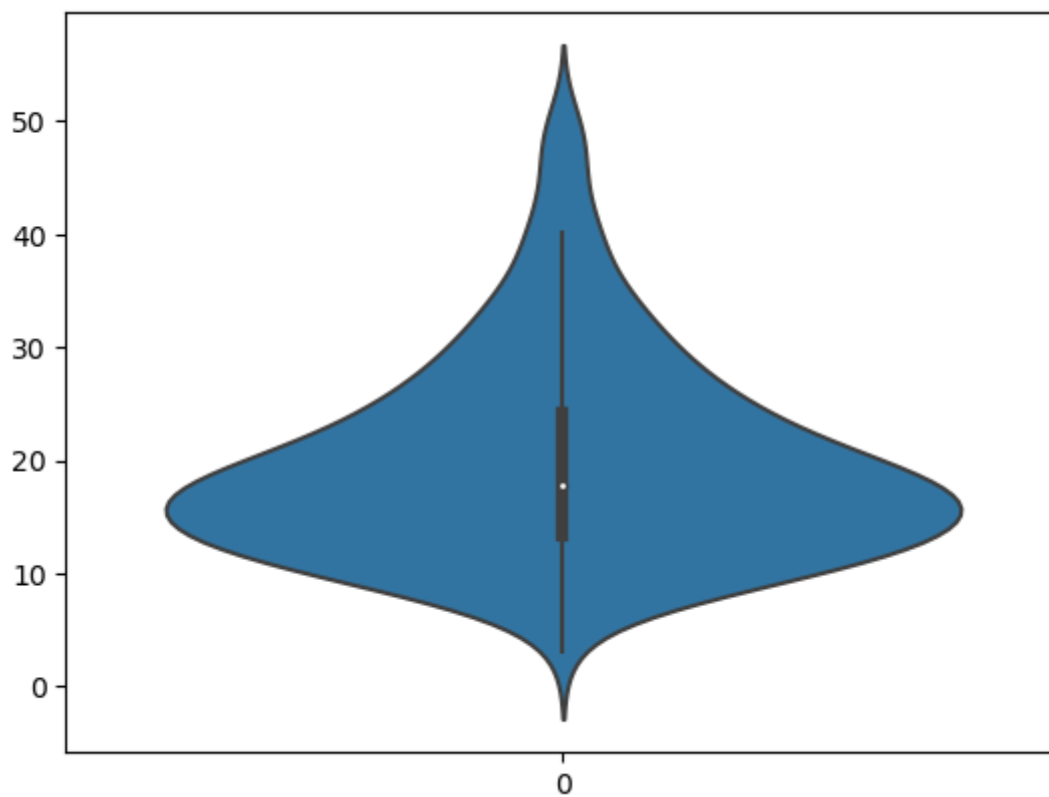


In [36]:

```
sn.violinplot(tips.total_bill)
```

Out[36]:

<Axes: >

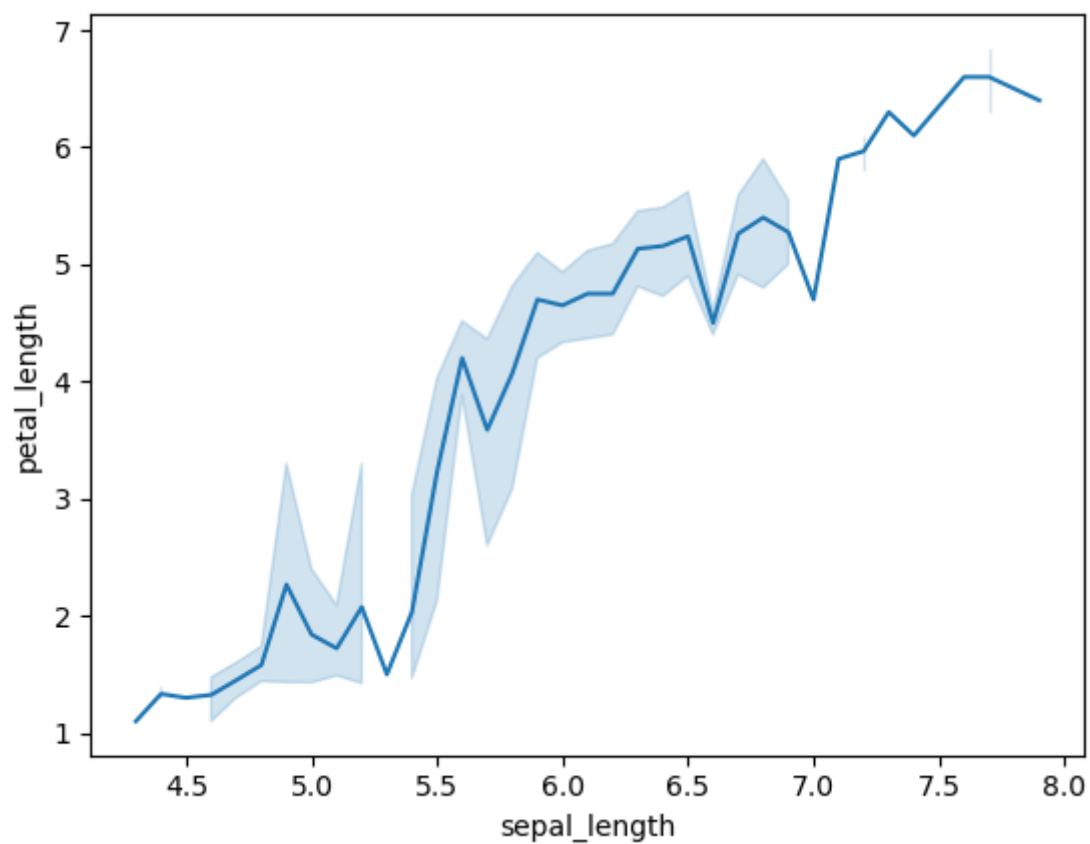


In [37]:

```
sn.lineplot(x= iris.sepal_length,y = iris.petal_length)
```

Out[37]:

<Axes: xlabel='sepal\_length', ylabel='petal\_length'>

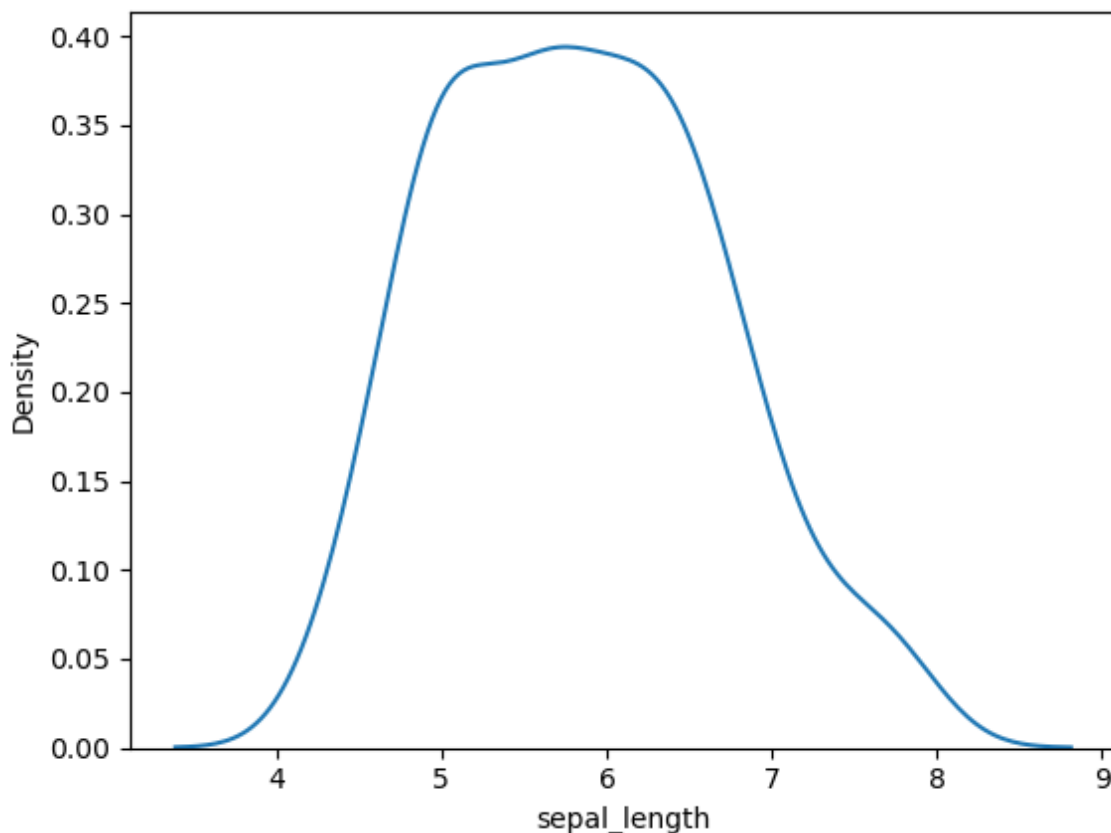


In [38]:

```
sn.kdeplot(iris.sepal_length)
```

Out[38]:

<Axes: xlabel='sepal\_length', ylabel='Density'>



**#heatmap** Heatmap is defined as a graphical representation of data using colors to visualize the value of the matrix. In this, to represent more common values or higher activities brighter colors basically reddish colors are used and to represent less common or activity values, darker colors are preferred. Heatmap is also defined by the name of the shading matrix. Heatmaps in Seaborn can be plotted by using the `seaborn.heatmap()` function.

`seaborn.heatmap()` Syntax: `seaborn.heatmap(data, *, vmin=None, vmax=None, cmap=None, center=None, annot_kws=None, linewidths=0, linecolor='white', cbar=True, **kwargs)`

**Important Parameters:**

**data:** 2D dataset that can be coerced into an ndarray. **vmin, vmax:** Values to anchor the colormap, otherwise they are inferred from the data and other keyword arguments. **cmap:** The mapping from data values to color space. **center:** The value at which to center the colormap when plotting divergent data. **annot:** If True, write the data value in each cell. **fmt:** String formatting code to use when adding annotations. **linewidths:** Width of the lines that will divide each cell. **linecolor:** Color of the lines that will divide each cell. **cbar:** Whether to draw a colorbar.

In [47]:

```
# importing the modules
import numpy as np
import seaborn as sn
import matplotlib.pyplot as plt

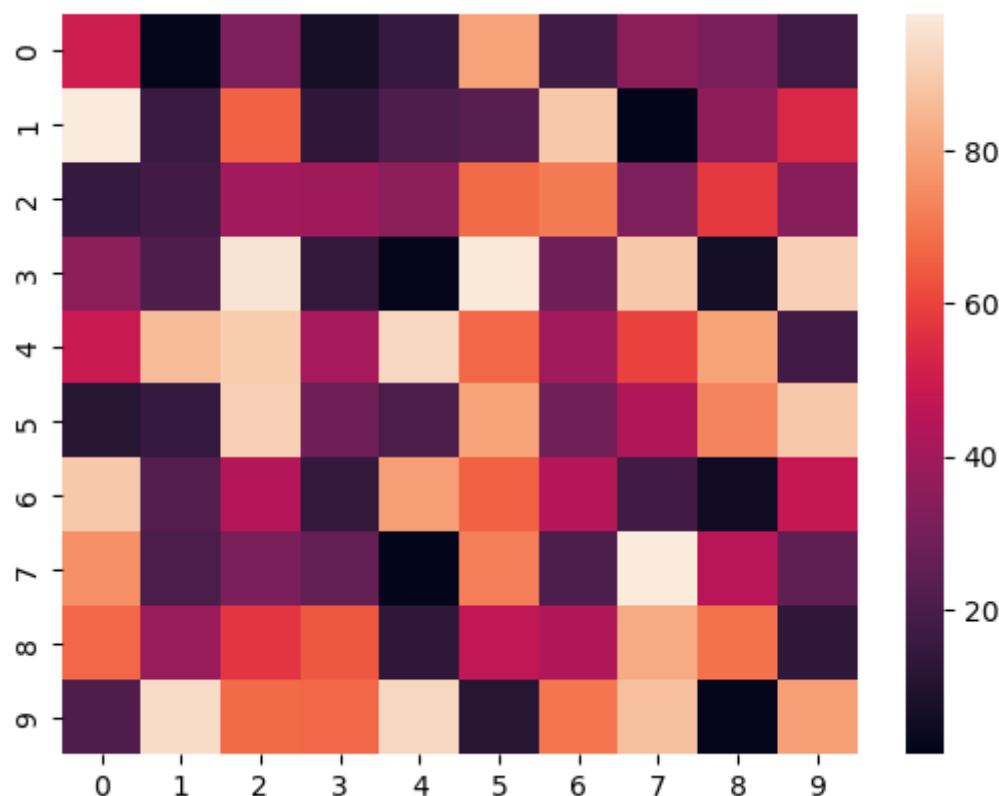
# generating 2-D 10x10 matrix of random numbers
# from 1 to 100
data = np.random.randint(low = 1,
                        high = 100,
                        size = (10, 10))

print("The data to be plotted:\n")
print(data)

# plotting the heatmap
hm = sn.heatmap(data = data)
```

The data to be plotted:

```
[[50  2 32  7 15 80 17 35 31 17]
 [98 16 66 13 21 23 89  1 36 54]
 [15 18 40 39 35 68 71 32 58 34]
 [35 21 96 14  2 97 28 89  6 91]
 [49 86 90 41 93 67 40 60 80 18]
 [11 15 91 28 20 80 29 43 73 89]
 [89 22 44 14 79 66 44 18  5 48]
 [76 20 31 26  1 72 20 98 45 25]
 [67 38 57 64 13 47 43 82 69 13]
 [21 94 68 67 93 11 70 87  2 79]]
```





In [51]:

```

import numpy as np
import seaborn as sn
import matplotlib.pyplot as plt

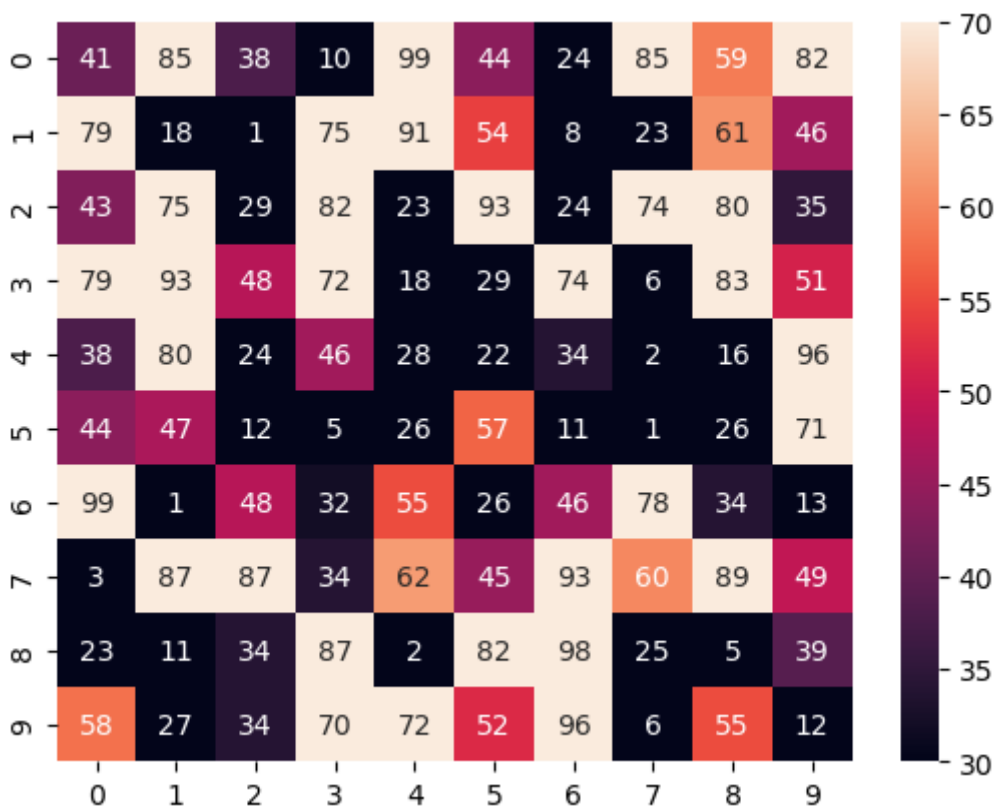
# generating 2-D 10x10 matrix of random numbers
# from 1 to 100
data = np.random.randint(low=1,
                        high=100,
                        size=(10, 10))

# setting the parameter values
vmin = 30
vmax = 70
#If we set the vmin value to 30 and the vmax value to 70, then only the cells with value

# plotting the heatmap
hm = sn.heatmap(data=data,
                vmin=vmin,
                vmax=vmax,
                annot= True
                )

# displaying the plotted heatmap
plt.show()
#If we set the vmin value to 30 and the vmax value to 70, then only the cells with value

```



In [53]:

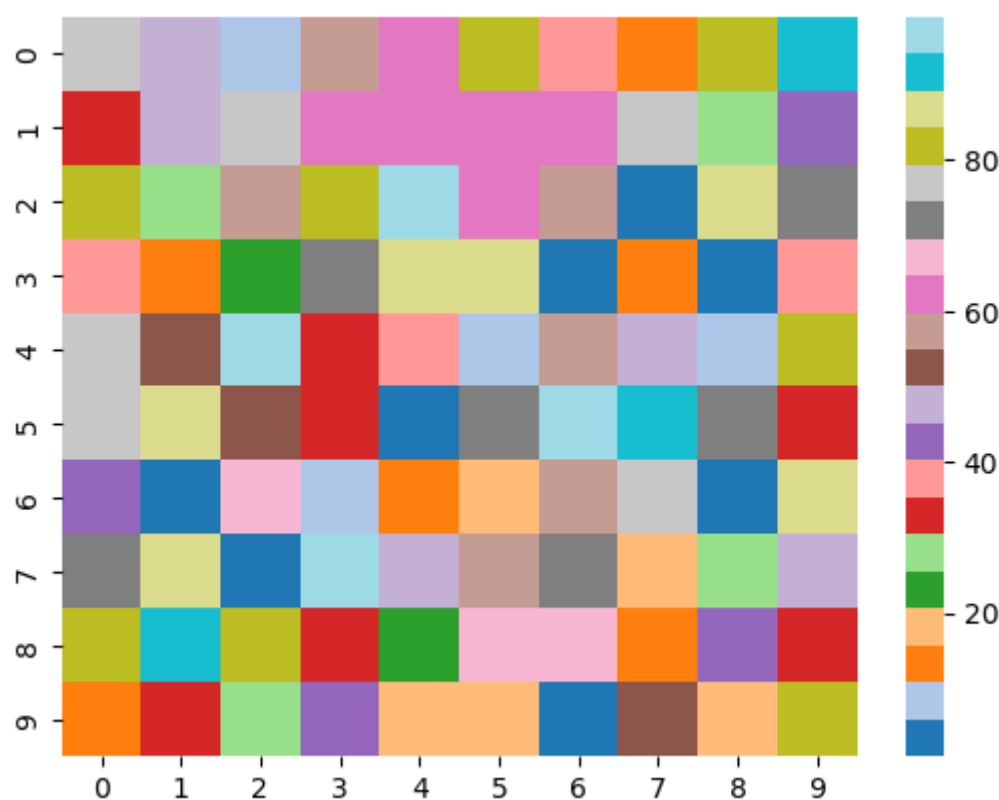
```
# importing the modules
import numpy as np
import seaborn as sn
import matplotlib.pyplot as plt

# generating 2-D 10x10 matrix of random numbers
# from 1 to 100
data = np.random.randint(low=1,
                        high=100,
                        size=(10, 10))

# setting the parameter values
cmap = "tab20"

# plotting the heatmap
hm = sn.heatmap(data=data,
                cmap=cmap)

# displaying the plotted heatmap
plt.show()
```



In [ ]: