

```
import matplotlib.pyplot as plt
import seaborn as sns
import warnings
warnings.filterwarnings('ignore')

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

Distribution Plots

- Distribution Plots are used for examining univariate and bivariate distributions meaning such distributions that involve one variable or two discrete variables.
- There are various types of categorical plots:
 - a. Histogram
 - b. Distplot
 - c. Pairplot
 - d. KDE Plot

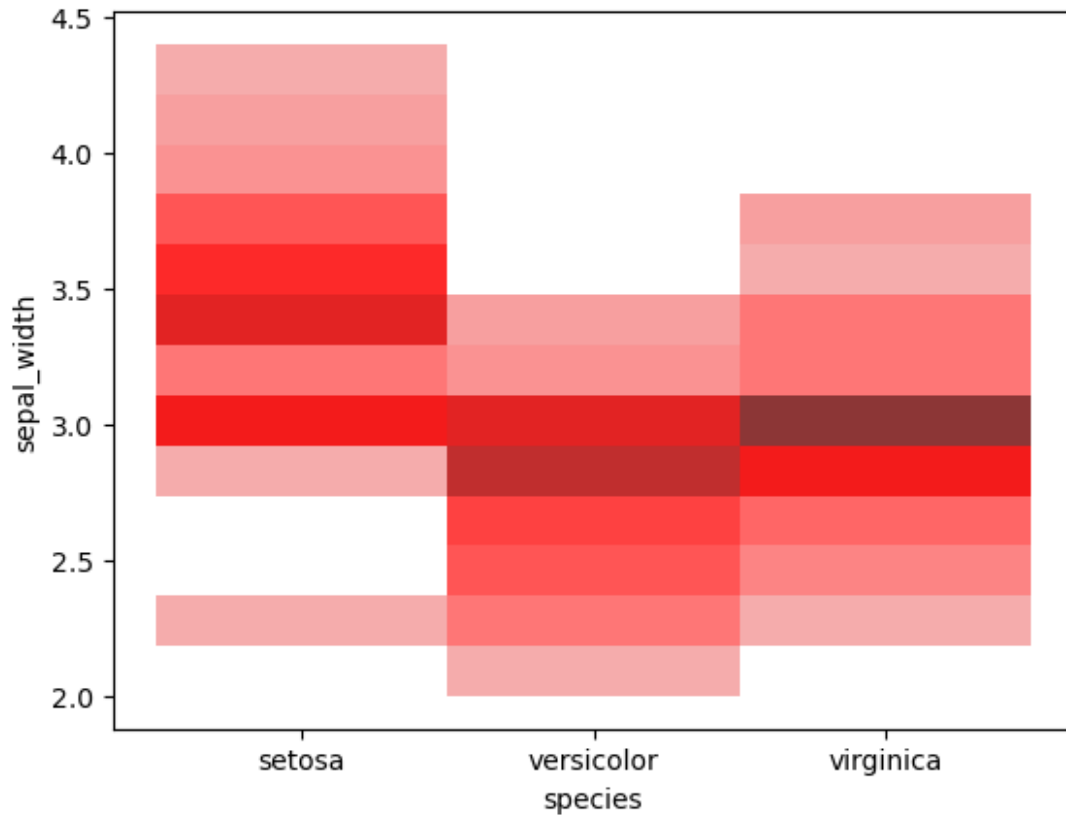
1. Histogram:

- A histogram is basically used to represent data provided in a form of some groups.
- It is accurate method for the graphical representation of numerical data distribution.
- It can be plotted using the `histplot()` function.
- Syntax:

```

- histplot(data=None, *, x=None, y=None, hue=None, **kwargs)
sns.histplot(x='species', y='sepal_width', data=iris, color='red')
plt.show()

```



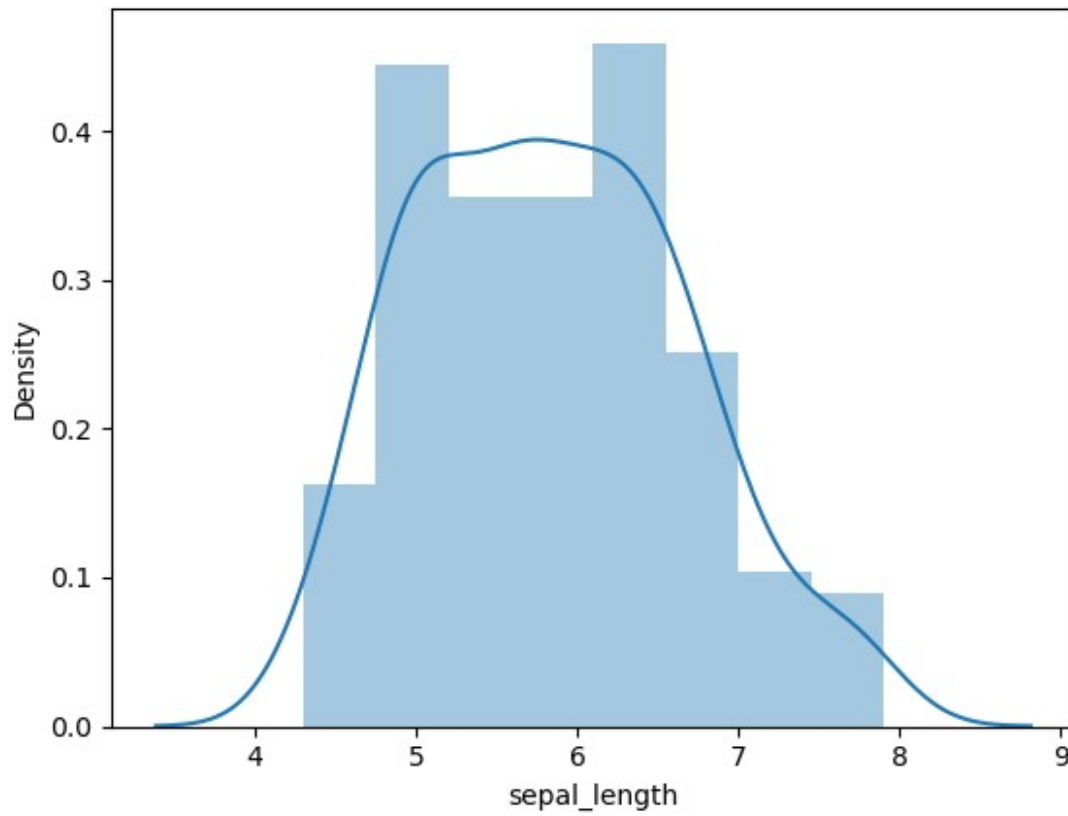
2. Distplot:

- Distplot is used basically for univariant set of observations and visualizes it through a histogram i.e. only one observation and hence we choose one particular column of the dataset.
- It is potted using the distplot() method.
- Syntax:

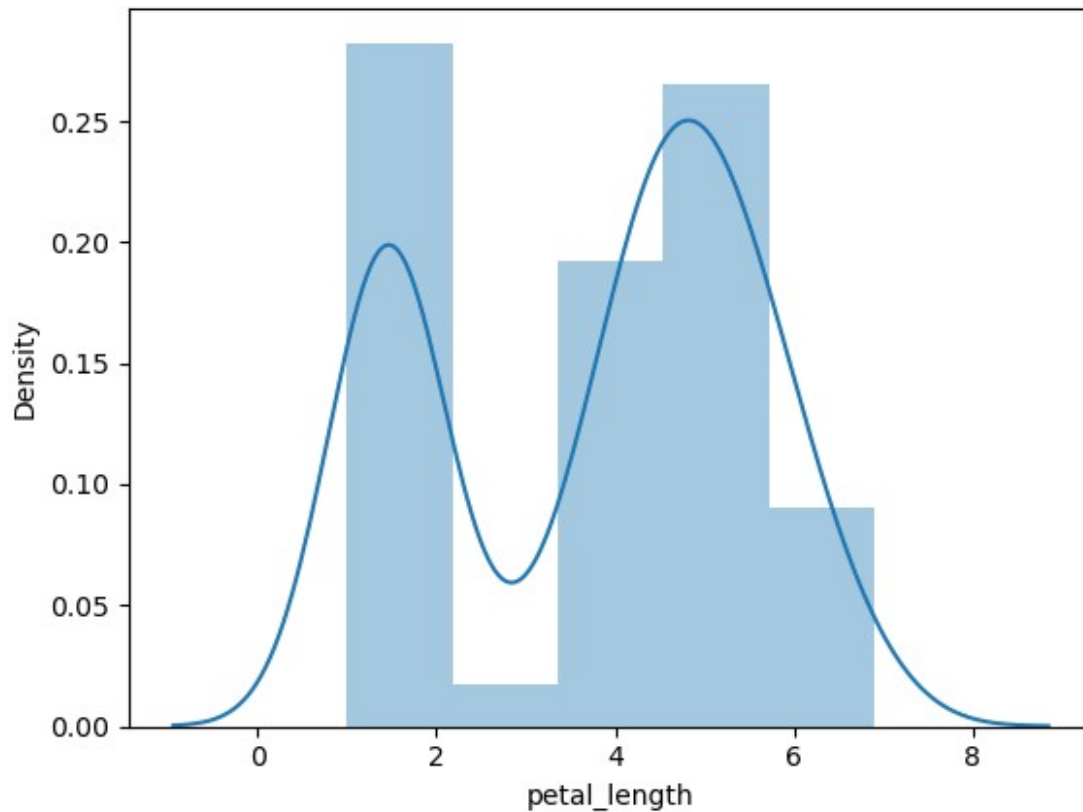
```

- distplot(a[, bins, hist, kde, rug, fit, ...])
sns.distplot(iris['sepal_length'])
plt.show()

```



```
sns.distplot(iris['petal_length'])  
plt.show()
```



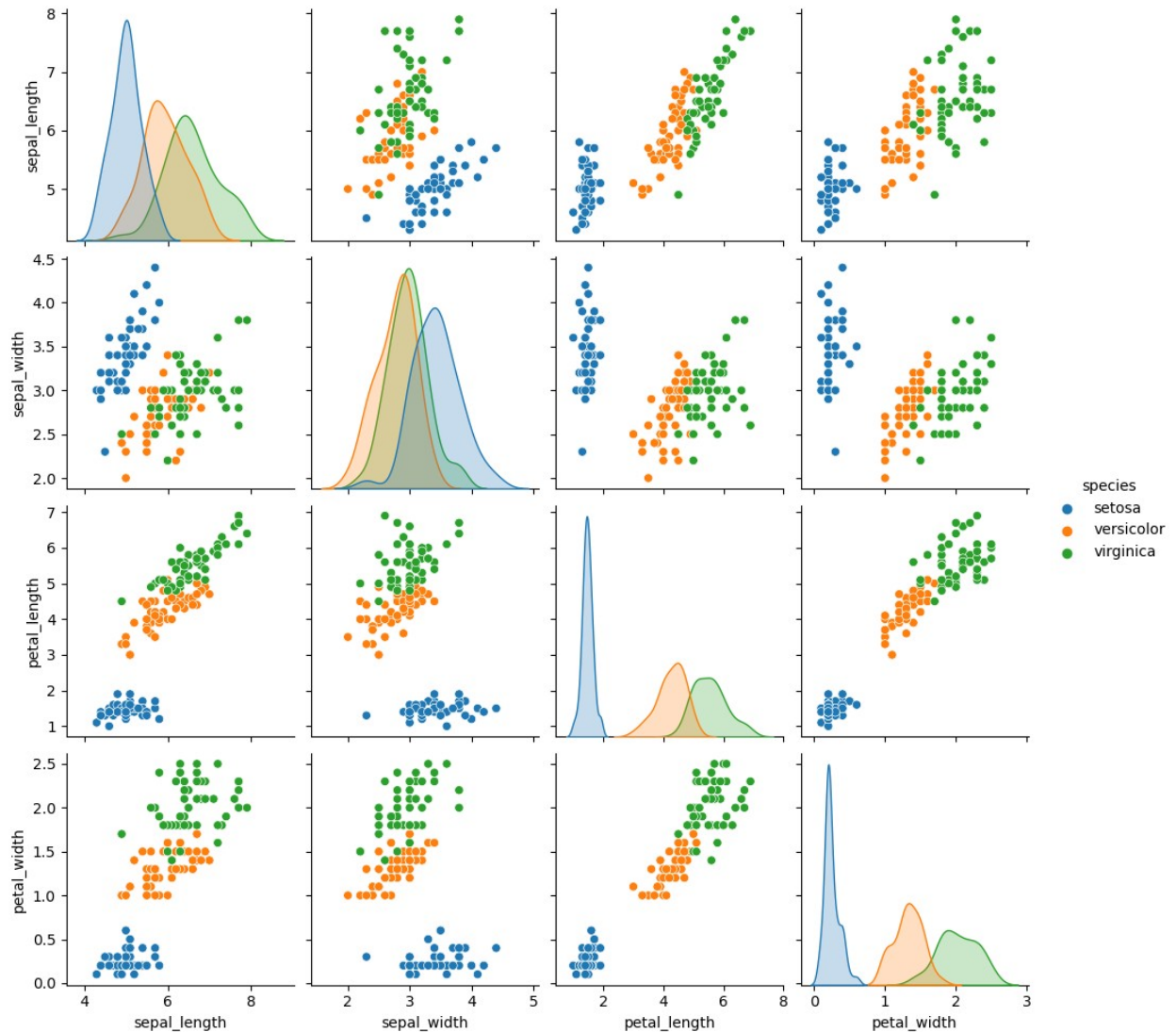
3. Pairplot:

- Pairplot represents pairwise relation across the entire dataframe and supports an additional argument called hue for categorical separation.
- It is plotted using the pairplot() method.
- Syntax:

– `pairplot(data[, hue, hue_order, palette, ...])`

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

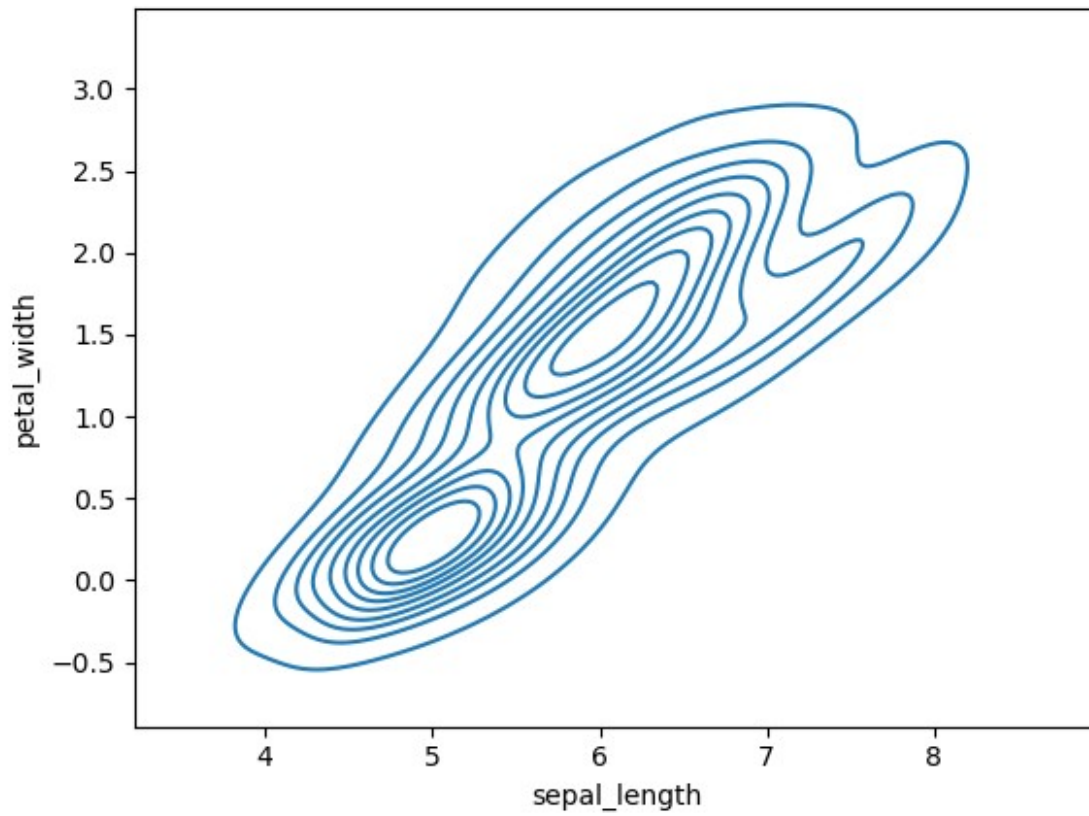
```
<seaborn.axisgrid.PairGrid at 0x1f8b8006510>
```



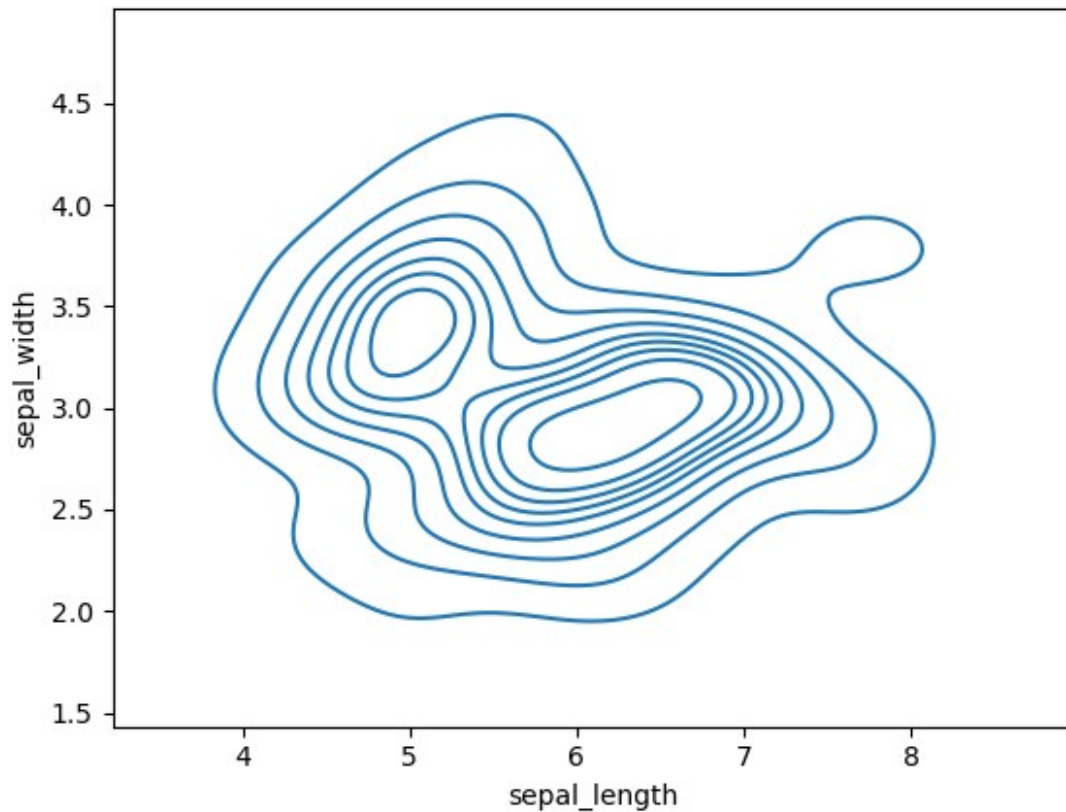
4. KDE Plot:

- KDE Plot described as Kernel Density Estimate is used for visualizing the Probability Density of a continuous variable.
- It depicts the probability density at different values in a continuous variable.
- We can also plot a single graph for multiple samples which helps in more efficient data visualization.
- Syntax:
 - `seaborn.kdeplot(x=None, *, y=None, vertical=False, palette=None, **kwargs)`

```
sns.kdeplot(x='sepal_length', y='petal_width', data=iris)
plt.show()
```



```
sns.kdeplot(x='sepal_length', y='sepal_width', data=data)  
plt.show()
```



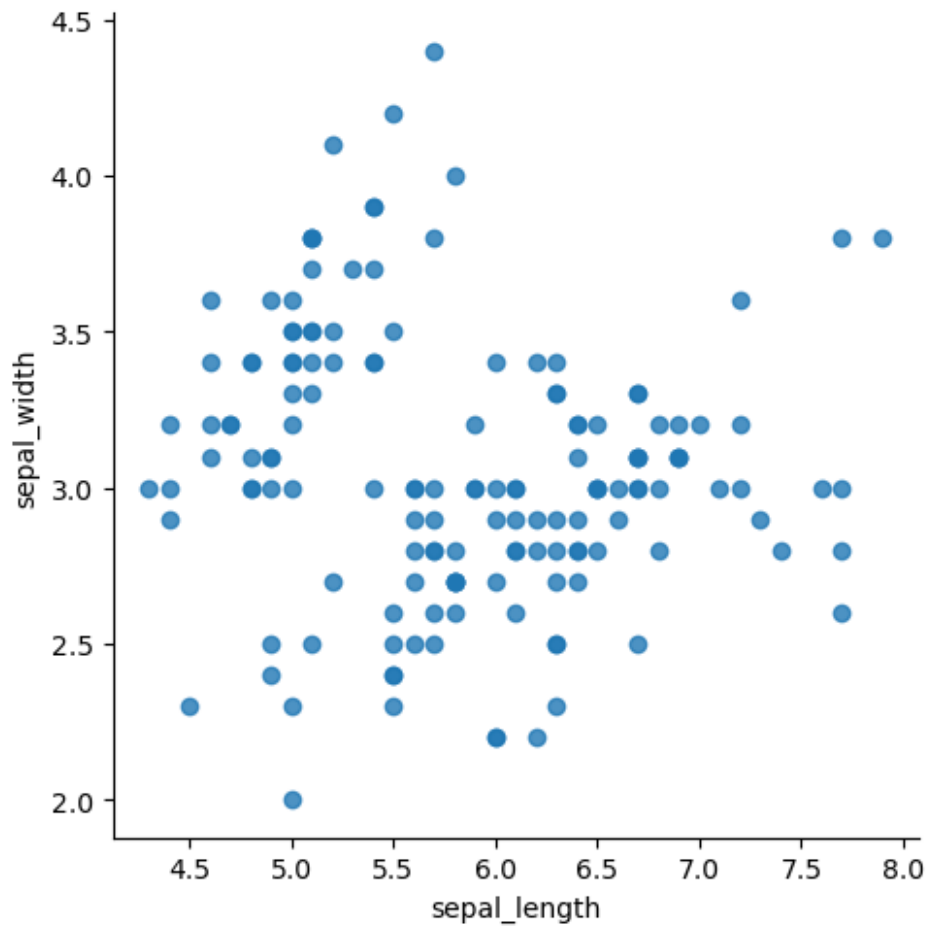
Regression Plots

- Regression plots as the name suggests creates a regression line between two parameters and helps to visualize their linear relationships.
- There are various types of categorical plots:
 - a. Implot
 - b. Regplot

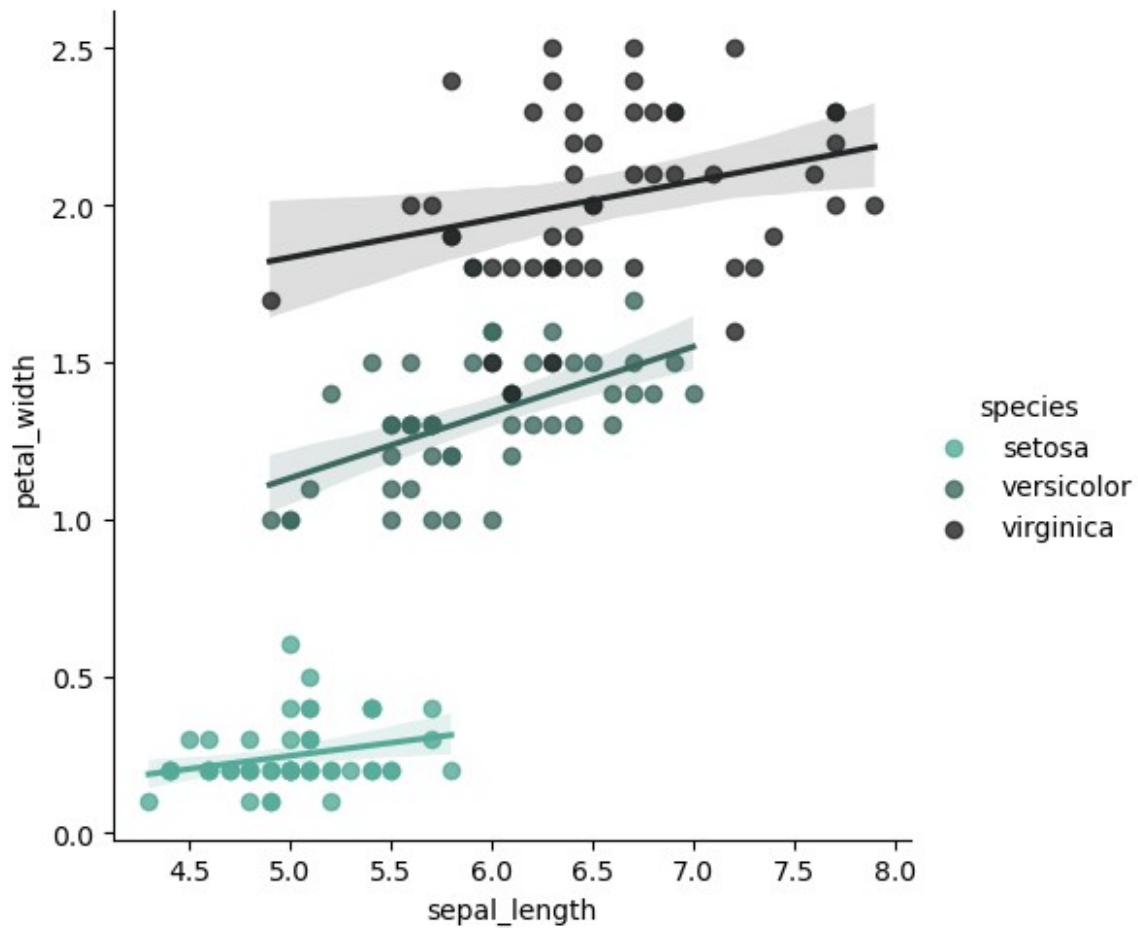
1. Implot:

- Implot() method can be understood as a function that basically creates a linear model plot.
- It creates a scatter plot with a linear fit on top of it.
- Syntax:
 - `seaborn.lmplot(x, y, data, hue=None, col=None, row=None, **kwargs)`

```
sns.lmplot(x='sepal_length', y='sepal_width', data=iris,  
fit_reg=False)  
plt.show()
```



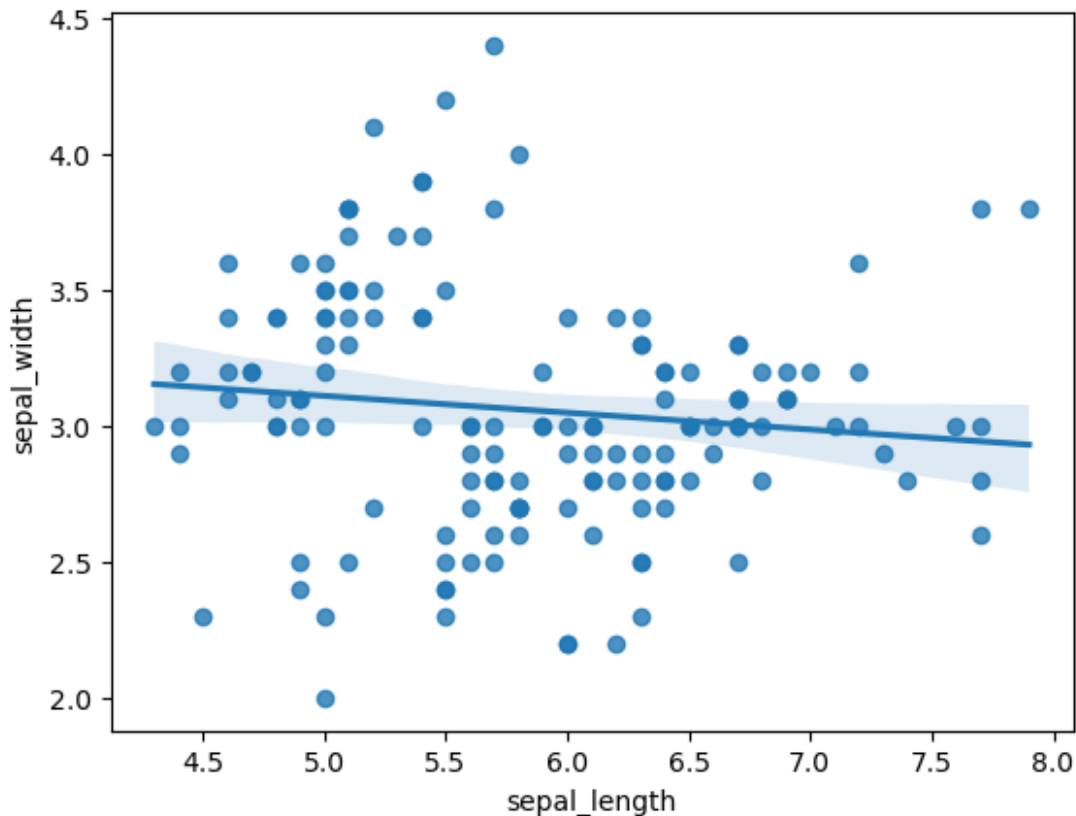
```
sns.lmplot(x='sepal_length', y='petal_width', data=iris,  
hue='species', palette='dark:#5A9_r')  
plt.show()
```

2. Regplot:

- regplot() method is also similar to lmpot which creates linear regression model.
- Syntax:
 - `seaborn.regplot(x, y, data=None, x_estimator=None, **kwargs)`

```
sns.regplot(x='sepal_length', y='sepal_width', data=iris)
plt.show()
```



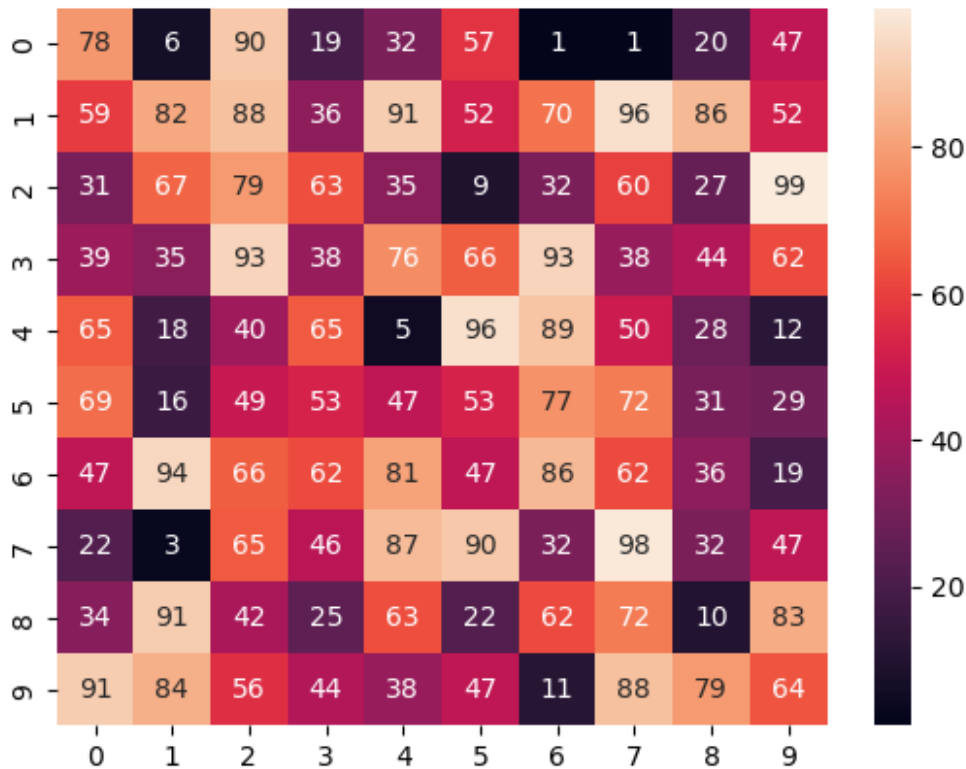
Matrix Plots

- A matrix plot means plotting matrix data, where color coded diagrams shows rows data, column data and values.
- It can shown using the heatmap and clustermap.
- There are various types of categorical plots:
 - a. Heatmap
 - b. Clustermap

1. 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.
- It can be plotted using the heatmap() function.
- Syntax:
 - `seaborn.heatmap(data, *, vmin=None, vmax=None, cmap=None, linecolor='white', cbar=True, **kwargs)`

```
import numpy as np
import pandas as pd
data = np.random.randint(low=1, high=100, size=(10, 10))
hm = sns.heatmap(data=data, annot=True)
plt.show()
```



2. Clustermmap:

- The `clustermmap()` function of seaborn plots the hierarchically-clustered heatmap of the given matrix dataset.
- Clustering simply means grouping data based on relationship among the variables in the data.
- Syntax:

```
– clustermmap(data, *, pivot_kws=None, **kwargs)
```

```
s1 = [100, 94, 56, 76, 81, 91, 51, 55, 72, 66, 60, 58 ]
s2 = [82, 81, 94, 96, 93, 84, 80, 82, 84, 86, 81, 78]
s3 = [65, 61, 66, 62, 67, 71, 69, 73, 68, 64, 66, 70]
s4 = [150, 140, 145, 151, 156, 152, 160, 165, 159, 149, 155, 162]
s5 = [75, 74, 76, 78, 80, 82, 85, 81, 77, 73, 75, 67]
s6 = [80, 75, 70, 72, 67, 65, 62, 63, 65, 60, 66, 69]
```

```
months= ["Jan", "Feb", "Mar", "Apr",
```

```

    "May", "Jun", "Jul", "Aug",
    "Sep", "Oct", "Nov", "Dec"]

d1 = {"State1":s1,
      "State2":s2,
      "State3":s3,
      "State4":s4,
      "State5":s5,
      "State6":s6};

df = pd.DataFrame(data=d1, index=months)
print(df)
print(df.columns)
print(df.index)

sns.clustermap(df)
plt.show()

```

	State1	State2	State3	State4	State5	State6
Jan	100	82	65	150	75	80
Feb	94	81	61	140	74	75
Mar	56	94	66	145	76	70
Apr	76	96	62	151	78	72
May	81	93	67	156	80	67
Jun	91	84	71	152	82	65
Jul	51	80	69	160	85	62
Aug	55	82	73	165	81	63
Sep	72	84	68	159	77	65
Oct	66	86	64	149	73	60
Nov	60	81	66	155	75	66
Dec	58	78	70	162	67	69

```

Index(['State1', 'State2', 'State3', 'State4', 'State5', 'State6'],
      dtype='object')
Index(['Jan', 'Feb', 'Mar', 'Apr', 'May', 'Jun', 'Jul', 'Aug', 'Sep',
      'Oct',
      'Nov', 'Dec'],
      dtype='object')

```

