

# Iris Flower:

## Importing all module

```
In [1]: 1 import pandas as pd
        2 import numpy as np
        3 import seaborn as sns
        4 import matplotlib.pyplot as plt
        5 import warnings
        6 warnings.filterwarnings('ignore')
```

## Load Datasets

```
In [2]: 1 #from sklearn.datasets import load_iris
        2 #iris=load_iris()
```

```
In [3]: 1 df=pd.read_csv('IRIS.csv')
```

```
In [4]: 1 df
```

```
Out[4]:
```

	sepal_length	sepal_width	petal_length	petal_width	species
0	5.1	3.5	1.4	0.2	Iris-setosa
1	4.9	3.0	1.4	0.2	Iris-setosa
2	4.7	3.2	1.3	0.2	Iris-setosa
3	4.6	3.1	1.5	0.2	Iris-setosa
4	5.0	3.6	1.4	0.2	Iris-setosa
...	...	...	...	...	...
145	6.7	3.0	5.2	2.3	Iris-virginica
146	6.3	2.5	5.0	1.9	Iris-virginica
147	6.5	3.0	5.2	2.0	Iris-virginica
148	6.2	3.4	5.4	2.3	Iris-virginica
149	5.9	3.0	5.1	1.8	Iris-virginica

150 rows × 5 columns

```
In [5]: 1 #iris.feature_names
        2 df.species
```

```
Out[5]: 0      Iris-setosa
        1      Iris-setosa
        2      Iris-setosa
        3      Iris-setosa
        4      Iris-setosa
        ...
       145      Iris-virginica
       146      Iris-virginica
       147      Iris-virginica
       148      Iris-virginica
       149      Iris-virginica
       Name: species, Length: 150, dtype: object
```

```
In [6]: 1 #df = pd.DataFrame(iris.data,columns=iris.feature_names)
```

```
In [7]: 1 #To display first five rows
        2 df.head()
```

```
Out[7]:
```

	sepal_length	sepal_width	petal_length	petal_width	species
0	5.1	3.5	1.4	0.2	Iris-setosa
1	4.9	3.0	1.4	0.2	Iris-setosa
2	4.7	3.2	1.3	0.2	Iris-setosa
3	4.6	3.1	1.5	0.2	Iris-setosa
4	5.0	3.6	1.4	0.2	Iris-setosa

```
In [8]: 1 #To display last five rows
        2 df.tail()
```

```
Out[8]:
```

	sepal_length	sepal_width	petal_length	petal_width	species
145	6.7	3.0	5.2	2.3	Iris-virginica
146	6.3	2.5	5.0	1.9	Iris-virginica
147	6.5	3.0	5.2	2.0	Iris-virginica
148	6.2	3.4	5.4	2.3	Iris-virginica
149	5.9	3.0	5.1	1.8	Iris-virginica

```
In [9]: 1 #df['class']=iris.target
        2 df.species.value_counts()
```

```
Out[9]: Iris-setosa      50
        Iris-versicolor  50
        Iris-virginica   50
        Name: species, dtype: int64
```

```
In [10]: 1 #to display datatypes
         2 df.dtypes
```

```
Out[10]: sepal_length    float64
          sepal_width    float64
          petal_length    float64
          petal_width    float64
          species         object
          dtype: object
```

```
In [11]: 1 df.shape
```

```
Out[11]: (150, 5)
```

```
In [12]: 1 df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 150 entries, 0 to 149
Data columns (total 5 columns):
#   Column          Non-Null Count  Dtype
---  ---
0   sepal_length    150 non-null   float64
1   sepal_width     150 non-null   float64
2   petal_length    150 non-null   float64
3   petal_width     150 non-null   float64
4   species         150 non-null   object
dtypes: float64(4), object(1)
memory usage: 6.0+ KB
```

```
In [13]: 1 #There are 150 observations with 4 features each (sepal length, sepal w
         2 #There are no null values, so we don't have to worry about that.
         3 #There are 50 observations of each species (setosa, versicolor, virgini
```

```
In [14]: 1 #to display the content of data
         2 df.describe()
```

```
Out[14]:
```

	sepal_length	sepal_width	petal_length	petal_width
count	150.000000	150.000000	150.000000	150.000000
mean	5.843333	3.054000	3.758667	1.198667
std	0.828066	0.433594	1.764420	0.763161
min	4.300000	2.000000	1.000000	0.100000
25%	5.100000	2.800000	1.600000	0.300000
50%	5.800000	3.000000	4.350000	1.300000
75%	6.400000	3.300000	5.100000	1.800000
max	7.900000	4.400000	6.900000	2.500000

```
In [15]: 1 # to display no. of samples of each Sepal Length
         2 df['sepal_length'].value_counts()
```

```
Out[15]: 5.0    10
         5.1     9
         6.3     9
         5.7     8
         6.7     8
         5.8     7
         5.5     7
         6.4     7
         4.9     6
         5.4     6
         6.1     6
         6.0     6
         5.6     6
         4.8     5
         6.5     5
         6.2     4
         7.7     4
         6.9     4
         4.6     4
         5.2     4
         5.9     3
         4.4     3
         7.2     3
         6.8     3
         6.6     2
         4.7     2
         7.6     1
         7.4     1
         7.3     1
         7.0     1
         7.1     1
         5.3     1
         4.3     1
         4.5     1
         7.9     1
         Name: sepal_length, dtype: int64
```

```
In [16]: 1 # to check for null values
         2 df.isnull().sum()
```

```
Out[16]: sepal_length    0
         sepal_width     0
         petal_length    0
         petal_width     0
         species         0
         dtype: int64
```

In [17]:

```
1 df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 150 entries, 0 to 149
Data columns (total 5 columns):
#   Column          Non-Null Count  Dtype  
---  -
0   sepal_length    150 non-null   float64
1   sepal_width     150 non-null   float64
2   petal_length    150 non-null   float64
3   petal_width     150 non-null   float64
4   species         150 non-null   object  
dtypes: float64(4), object(1)
memory usage: 6.0+ KB
```

In [18]:

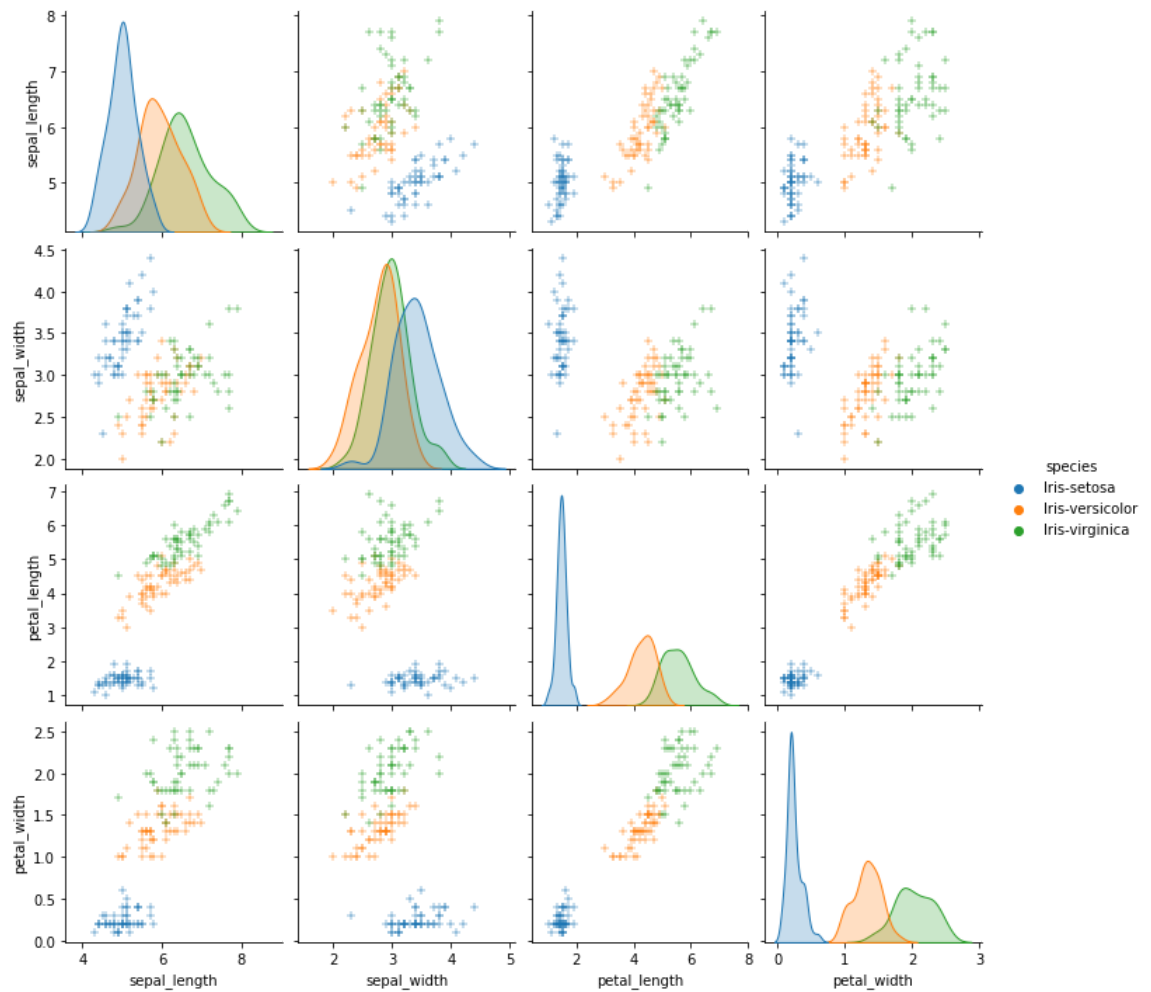
```
1 colname= df.select_dtypes('float64').columns
2 colname
```

Out[18]: Index(['sepal\_length', 'sepal\_width', 'petal\_length', 'petal\_width'], dtype='object')

## Data Visulazation

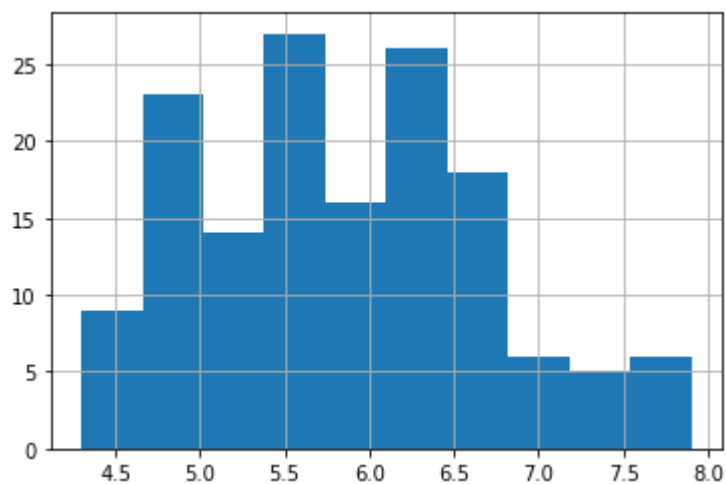
**After plotting the features in a pair plot, it is obvious that the connection between pairs of traits in an iris-setosa (in pink) differs significantly from those in the other two class. The paired relationships of the other two class, iris-versicolor (brown) and iris-virginica (green), have some overlap.**

```
In [19]: 1 g = sns.pairplot(df, hue='species', markers='+')
2         plt.show()
```



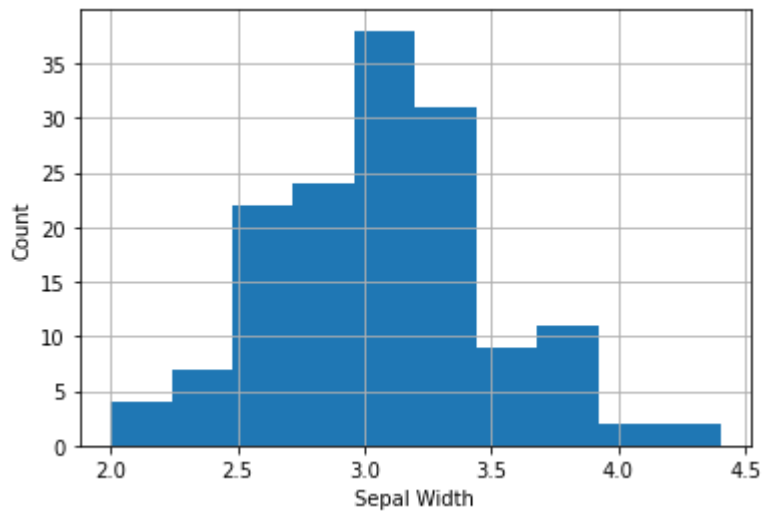
```
In [20]: 1 #Histogram
2         df["sepal_length"].hist()
```

Out[20]: <AxesSubplot:>



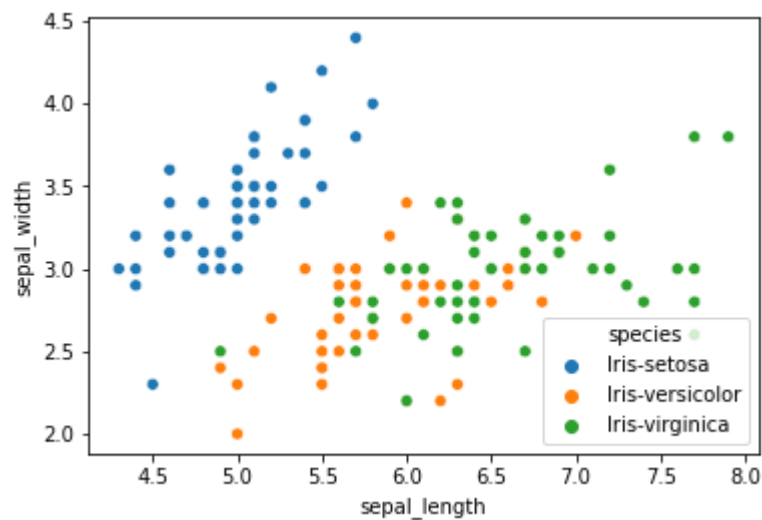
```
In [21]: 1 a = df["sepal_width"].hist()  
2 a.set_xlabel ("Sepal Width")  
3 a.set_ylabel ("Count")
```

Out[21]: Text(0, 0.5, 'Count')



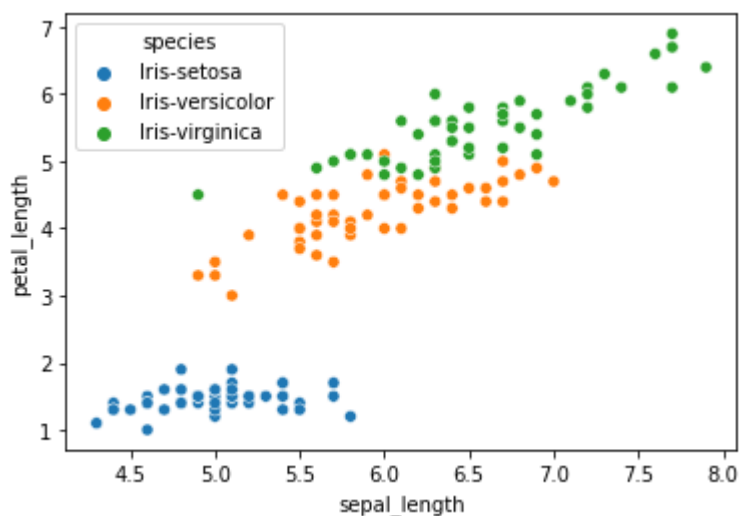
```
In [22]: 1 # Plotting Scatterplot using Seaborn  
2 sns.scatterplot(data=df, x='sepal_length', y='sepal_width', hue='species')
```

Out[22]: <AxesSubplot:xlabel='sepal\_length', ylabel='sepal\_width'>



```
In [23]: 1 # Plotting Scatterplot using Seaborn
          2 sns.scatterplot(data=df, x='sepal_length', y='petal_length', hue='species')
```

Out[23]: <AxesSubplot:xlabel='sepal\_length', ylabel='petal\_length'>



## Matrix of correlation

A correlation matrix is a table that contains the coefficients of correlation between different features (attributes) in a dataset. The correlation between two variables is represented by each cell in the table. The value ranges from -1 to 1.

```
In [24]: 1 # to delete Species Class from the dataset
          2 new_df = df.drop(columns=['species'])
          3 new_df.head()
```

Out[24]:

	sepal_length	sepal_width	petal_length	petal_width
0	5.1	3.5	1.4	0.2
1	4.9	3.0	1.4	0.2
2	4.7	3.2	1.3	0.2
3	4.6	3.1	1.5	0.2
4	5.0	3.6	1.4	0.2

```
In [25]: 1 # correlation matrix
          2 corr_mat = new_df.corr()
          3 corr_mat
```

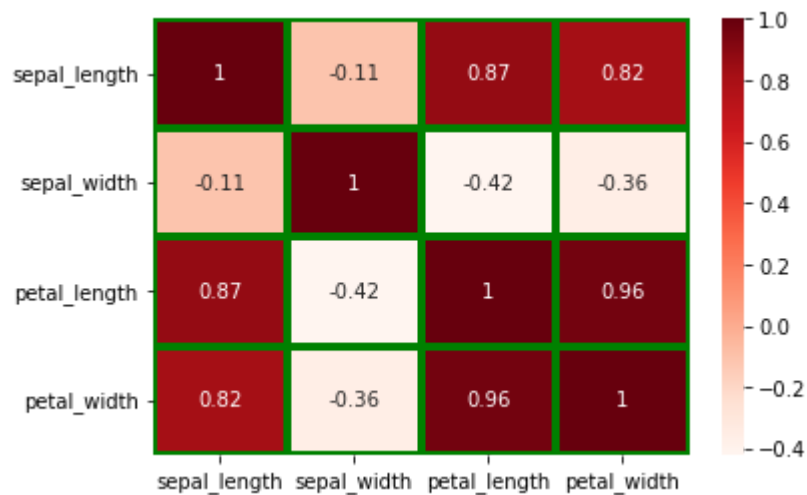
Out[25]:

	sepal_length	sepal_width	petal_length	petal_width
sepal_length	1.000000	-0.109369	0.871754	0.817954
sepal_width	-0.109369	1.000000	-0.420516	-0.356544
petal_length	0.871754	-0.420516	1.000000	0.962757
petal_width	0.817954	-0.356544	0.962757	1.000000



```
In [26]: 1 # Heatmap
          2 sns.heatmap(corr_mat, cmap='Reds', annot=True, linewidths=4, linecolor=
```

Out[26]: <AxesSubplot:>



```
In [27]: 1 #Label Encoder
```

```
In [28]: 1 #from scipy.stats import skew
```

```
In [29]: 1 #x=df.iloc[:, :-1]
          2 #y=df.iloc[:, -1]
          3 x = df.drop(columns='species')
          4 y = df['species']
```

```
In [30]: 1 x
```

```
Out[30]:
```

	sepal_length	sepal_width	petal_length	petal_width
0	5.1	3.5	1.4	0.2
1	4.9	3.0	1.4	0.2
2	4.7	3.2	1.3	0.2
3	4.6	3.1	1.5	0.2
4	5.0	3.6	1.4	0.2
...	...	...	...	...
145	6.7	3.0	5.2	2.3
146	6.3	2.5	5.0	1.9
147	6.5	3.0	5.2	2.0
148	6.2	3.4	5.4	2.3
149	5.9	3.0	5.1	1.8

150 rows × 4 columns

In [31]:

```
1 y
```

Out[31]:

```
0      Iris-setosa
1      Iris-setosa
2      Iris-setosa
3      Iris-setosa
4      Iris-setosa
...
145    Iris-virginica
146    Iris-virginica
147    Iris-virginica
148    Iris-virginica
149    Iris-virginica
Name: species, Length: 150, dtype: object
```

In [32]:

```
1 #Split the data train and test
```

In [33]:

```
1 from sklearn.model_selection import train_test_split
2 xtrain,xtest,ytrain,ytest=train_test_split(x,y,test_size=0.2,random_sta
```

In [34]:

```
1 #By building a model
2 def mymodel(model):
3
4     model.fit(xtrain,ytrain)
5     ypred= model.predict(xtest)
6
7     train=model.score(xtrain,ytrain)
8     test=model.score(xtest,ytest)
9
10    print(f'Training Accuracy:- {train}\nTesting Accuracy:- {test}')
11
12    print(classification_report(ytest,ypred))
13    return model
```

## Build Model

In [35]:

```
1 from sklearn.neighbors import KNeighborsClassifier
2 from sklearn.linear_model import LogisticRegression
3 from sklearn.svm import SVC
4 from sklearn.tree import DecisionTreeClassifier
```

In [36]:

```
1 from sklearn.metrics import classification_report
```

```
In [37]: 1 knn= mymodel(KNeighborsClassifier())
```

Training Accuracy:- 0.95

Testing Accuracy:- 0.9666666666666667

	precision	recall	f1-score	support
Iris-setosa	1.00	1.00	1.00	11
Iris-versicolor	1.00	0.92	0.96	13
Iris-virginica	0.86	1.00	0.92	6
accuracy			0.97	30
macro avg	0.95	0.97	0.96	30
weighted avg	0.97	0.97	0.97	30

```
In [38]: 1 logreg= mymodel(LogisticRegression())
```

Training Accuracy:- 0.9666666666666667

Testing Accuracy:- 1.0

	precision	recall	f1-score	support
Iris-setosa	1.00	1.00	1.00	11
Iris-versicolor	1.00	1.00	1.00	13
Iris-virginica	1.00	1.00	1.00	6
accuracy			1.00	30
macro avg	1.00	1.00	1.00	30
weighted avg	1.00	1.00	1.00	30

```
In [39]: 1 svm= mymodel(SVC())
```

Training Accuracy:- 0.9583333333333334

Testing Accuracy:- 1.0

	precision	recall	f1-score	support
Iris-setosa	1.00	1.00	1.00	11
Iris-versicolor	1.00	1.00	1.00	13
Iris-virginica	1.00	1.00	1.00	6
accuracy			1.00	30
macro avg	1.00	1.00	1.00	30
weighted avg	1.00	1.00	1.00	30

```
In [40]: 1 dt= mymodel(DecisionTreeClassifier())
```

Training Accuracy:- 1.0

Testing Accuracy:- 1.0

	precision	recall	f1-score	support
Iris-setosa	1.00	1.00	1.00	11
Iris-versicolor	1.00	1.00	1.00	13
Iris-virginica	1.00	1.00	1.00	6
accuracy			1.00	30
macro avg	1.00	1.00	1.00	30
weighted avg	1.00	1.00	1.00	30