

IRIS FLOWER CLASSIFICATION

Iris flower has three species: **setosa**, **versicolor**, and **virginica**

Which differs according to their measurements. Now assume that you have the measurements of the iris flowers according to their species, and here your task is to train a machine learning model that can learn from the measurements of the iris species and classify them.

Although the Scikit-learn library provides a dataset for iris flower classification, you can also download the same dataset from here for the task of iris flower classification with Machine Learning.

Import the required modules

```
In [51]: import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
import os
%matplotlib inline

from sklearn import metrics
from sklearn.neighbors import KNeighborsClassifier
from sklearn.linear_model import LogisticRegression
from sklearn.model_selection import train_test_split
```

Loading the IRIS dataset

```
In [14]: data = pd.read_csv("IRIS.csv")
data.head()
```

```
Out[14]:   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 [15]: data.tail()
```

```
Out[15]:   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 [16]: #displaying the statistics of data
data.describe()
```

```
Out[16]:   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 [17]: #basic info about datatype
data.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: 6.8+ KB
```

```
In [20]: #info about columns of data
data.columns
```

```
Out[20]: Index(['sepal_length', 'sepal_width', 'petal_length', 'petal_width',
              'species'],
              dtype='object')
```

```
In [22]: #to know no. of samples in each class(species variants)
data['species'].value_counts()
```

```
Out[22]: Iris-setosa      50
Iris-versicolor      50
Iris-virginica       50
Name: species, dtype: int64
```

Data Preprocessing

```
In [23]: #to know any null values in dataset
data.isnull().sum()

#glad that i dont't have any null values
```

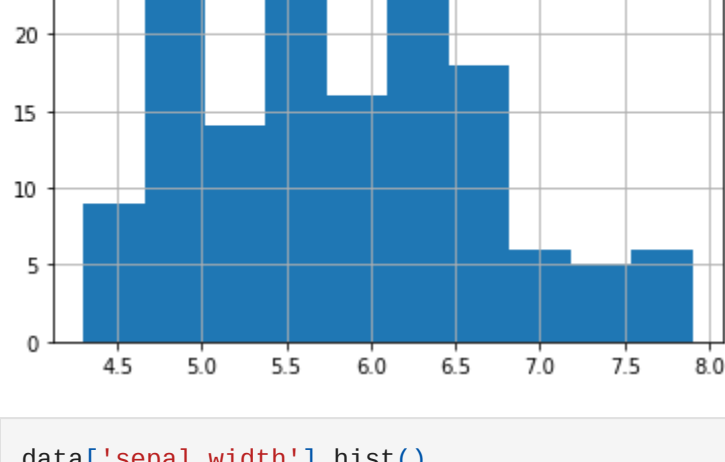
```
Out[23]: sepal_length      0
sepal_width      0
petal_length      0
petal_width      0
species          0
dtype: int64
```

Exploratory Data Analysis

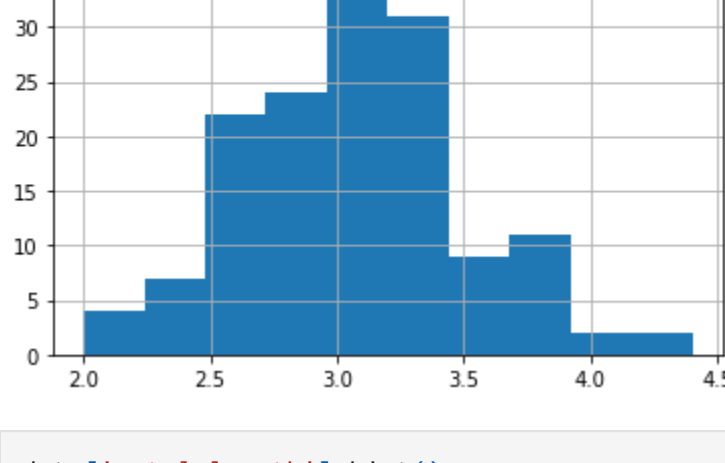
Here, I'll visualize the data in the form of graphs.

1. Histogram

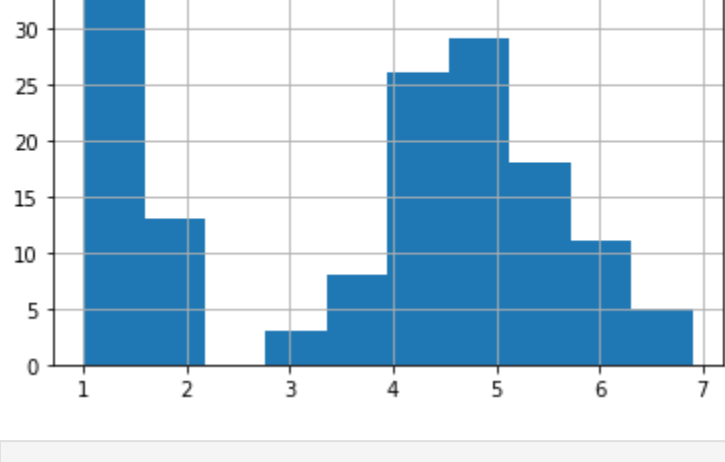
```
In [26]: data['sepal_length'].hist()
plt.show()
```



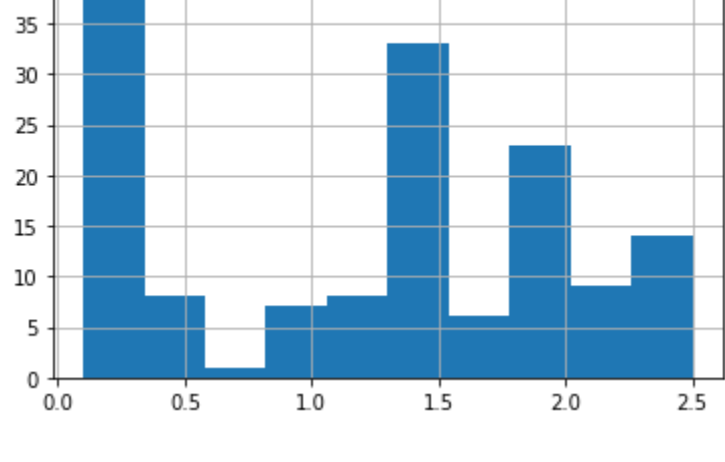
```
In [27]: data['sepal_width'].hist()
plt.show()
```



```
In [28]: data['petal_length'].hist()
plt.show()
```



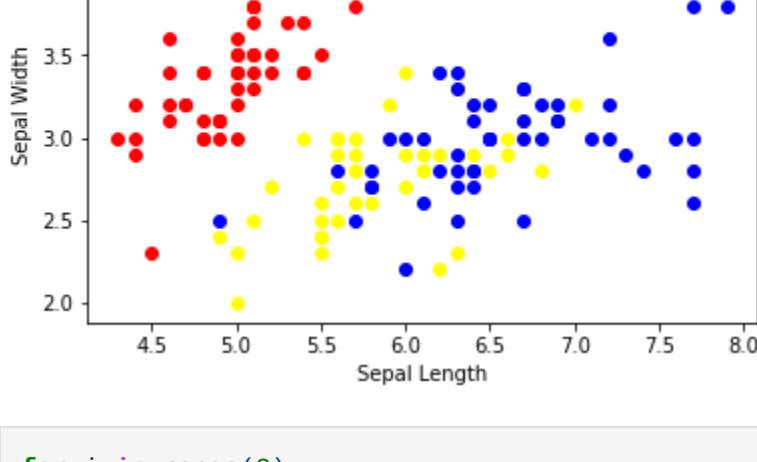
```
In [29]: data['petal_width'].hist()
plt.show()
```



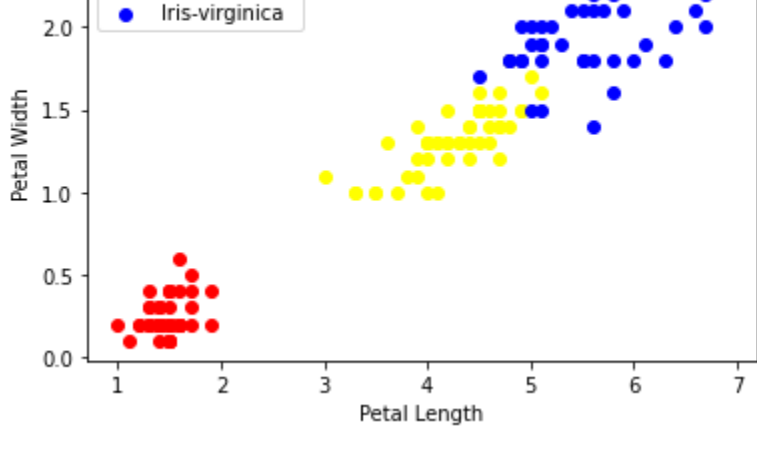
1. Scatterplot

```
In [36]: colors = ['red', 'yellow', 'blue']
species = ['Iris-setosa', 'Iris-versicolor', 'Iris-virginica']
```

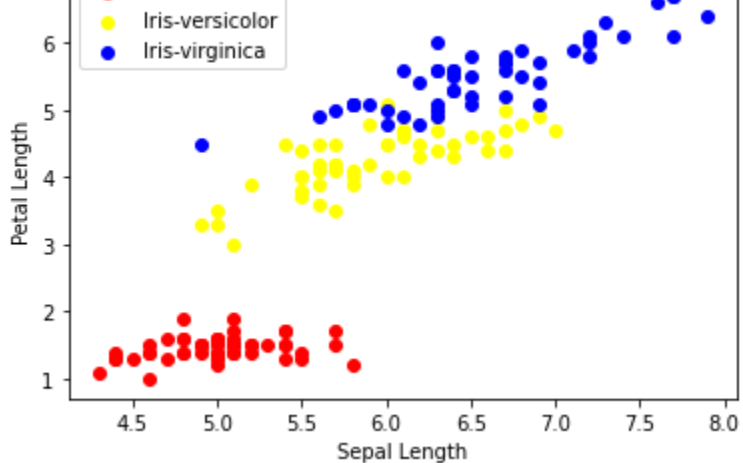
```
In [39]: for i in range(3):
x = data[data['species'] == species[i]]
plt.scatter(x['sepal_length'], x['sepal_width'], c = colors[i], label = species[i])
plt.xlabel("sepal_length")
plt.ylabel("sepal_width")
plt.legend()
plt.show()
```



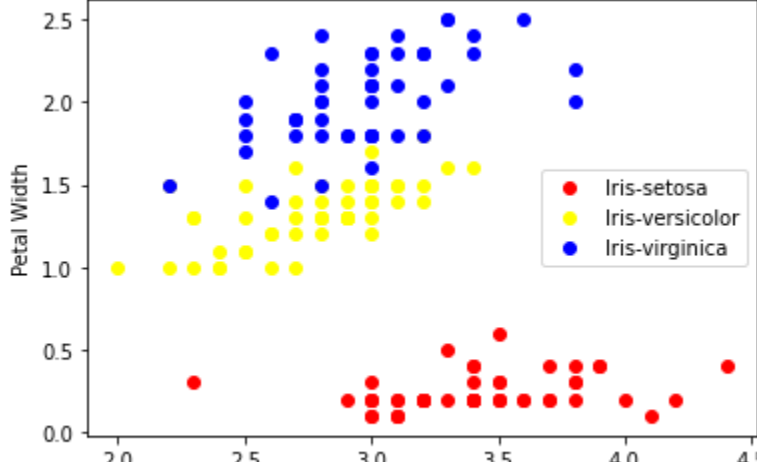
```
In [41]: for i in range(3):
x = data[data['species'] == species[i]]
plt.scatter(x['sepal_length'], x['petal_width'], c = colors[i], label = species[i])
plt.xlabel("sepal_length")
plt.ylabel("petal_width")
plt.legend()
plt.show()
```



```
In [42]: for i in range(3):
x = data[data['species'] == species[i]]
plt.scatter(x['sepal_length'], x['petal_length'], c = colors[i], label = species[i])
plt.xlabel("sepal_length")
plt.ylabel("petal_length")
plt.legend()
plt.show()
```



```
In [43]: for i in range(3):
x = data[data['species'] == species[i]]
plt.scatter(x['sepal_width'], x['petal_width'], c = colors[i], label = species[i])
plt.xlabel("sepal_width")
plt.ylabel("petal_width")
plt.legend()
plt.show()
```



Coorelation Matrix

Coorelation matrix is a table which shows the coorelation coefficients between variables.

Each cell in the table shows the correlation between variables.

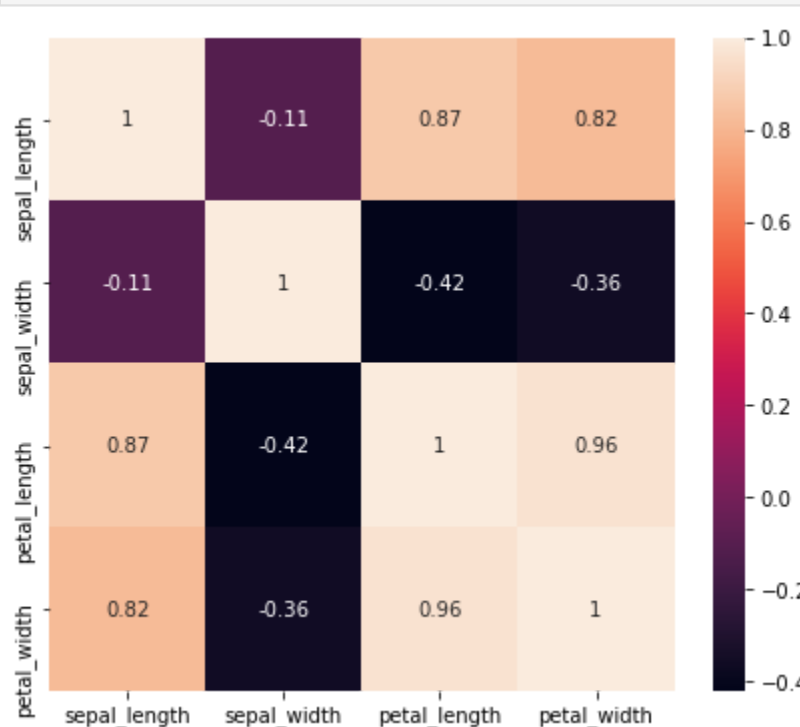
Value range of -1 to +1.

If two variables have high correlation, can neglect one of those.

```
In [45]: coor = data.corr()
coor
```

```
Out[45]:   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 [56]: fig, ax = plt.subplots(figsize =(7,6))
sns.heatmap(coor, annot = True, ax = ax)
plt.show()
```



Label Encoder

converts Categorical into Numeric label

```
In [59]: from sklearn.preprocessing import LabelEncoder
lab = LabelEncoder()
```

Iris-setosa as 0

Iris-versicolor as 1

Iris-virginica as 2

Name: species, dtype: int64

```
In [63]: data['species'] = lab.fit_transform(data['species'])
data.head()
```

```
Out[63]:   sepal_length  sepal_width  petal_length  petal_width  species
0          5.1           3.5           1.4           0.2      0
1          4.9           3.0           1.4           0.2      0
2          4.7           3.2           1.3           0.2      0
3          4.6           3.1           1.5           0.2      0
4          5.0           3.6           1.4           0.2      0
```

```
In [64]: data.tail()
```

```
Out[64]:   sepal_length  sepal_width  petal_length  petal_width  species
145          6.7           3.0           5.2           2.3      2
146          6.3           2.5           5.0           1.9      2
147          6.5           3.0           5.2           2.0      2
148          6.2           3.4           5.4           2.3      2
149          5.9           3.0           5.1           1.8      2
```

```
In [68]: data.columns
```

```
Out[68]: Index(['sepal_length', 'sepal_width', 'petal_length', 'petal_width',
              'species'],
              dtype='object')
```

```
In [85]: from sklearn.model_selection import train_test_split
#test = 30 and train = 70
x = data.drop(columns=['species'])
y = data['species']
x_train, x_test, y_train, y_test = train_test_split(X, y, test_size = 0.30)
```

```
In [86]: #logistic regression
from sklearn.linear_model import LogisticRegression
model = LogisticRegression()
```

```
In [87]: #model training
model.fit(x_train, y_train)
```

```
C:\ProgramData\Anaconda3\lib\site-packages\sklearn\linear_model\logistic.py:763: ConvergenceWarning: lbfgs failed to converge (status=1):
STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.

Increase the number of iterations (max_iter) or scale the data as shown in:
https://scikit-learn.org/stable/modules/preprocessing.html
Please also refer to the documentation for alternative solver options:
https://scikit-learn.org/stable/modules/linear_model.html#logistic-regression
n_iter_1 = check_optimize_result(
LogisticRegression()
```

```
In [123]: #printing the metric to know the performance of the model trained by LogisticRegression()
print('Accuracy of the model:', model.score(x_test, y_test) * 100)
```

Accuracy of the model: 84.44444444444444

```
In [95]: #knn - k nearest neighbours
from sklearn.neighbors import KNeighborsClassifier
model = KNeighborsClassifier()
```

```
In [97]: model.fit(x_train, y_train)
```

```
Out[97]: KNeighborsClassifier()
```

```
In [117]: #printing the metric to know the performance of the model trained by KNeighborsClassifier()
print('Accuracy of the model:', model.score(x_test, y_test) * 100)
```

Accuracy of the model: 84.44444444444444

```
In [109]: from sklearn import tree
```

```
In [114]: #decision tree
from sklearn.tree import DecisionTreeClassifier
model = DecisionTreeClassifier()
```

```
In [115]: model.fit(x_train, y_train)
```

```
Out[115]: DecisionTreeClassifier()
```

```
In [125]: #printing the metric to know the performance of the model trained by DecisionTreeClassifier()
print('Accuracy of the model:', model.score(x_test, y_test) * 100)
```

Accuracy of the model: 84.44444444444444