

## IMPORT LIBRARIES

```
In [3]: import pandas as pd
import numpy as np
import seaborn as sns
from matplotlib import pyplot as plt
import warnings
warnings.filterwarnings("ignore")
```

## LOAD DATASET

```
In [4]: df = pd.read_csv("IRIS.csv")
```

```
In [5]: df.head()
```

```
Out[5]:
```

|   | 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 [6]: 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
```

# MISSING VALUES

```
In [7]: print("sepal_length:")
print("Null values:",df['sepal_length'].isna().sum())
print("value counts:\n",df['sepal_length'].value_counts())
```

```
sepal_length:
Null values: 0
value counts:
```

```
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 [8]: print("sepal_width:")
print("Null_values:",df['sepal_width'].isna().sum())
print("value counts:\n",df['sepal_width'].value_counts())
```

```
sepal_width:
Null_values: 0
value counts:
 3.0    26
 2.8    14
 3.2    13
 3.1    12
 3.4    12
 2.9    10
 2.7     9
 2.5     8
 3.5     6
 3.3     6
 3.8     6
 2.6     5
 2.3     4
 3.7     3
 2.4     3
 2.2     3
 3.6     3
 3.9     2
 4.4     1
 4.0     1
 4.1     1
 4.2     1
 2.0     1
Name: sepal_width, dtype: int64
```

---

```
In [9]: print("petal_length:")
print("Null_values:",df['petal_length'].isna().sum())
print("value counts:\n",df['petal_length'].value_counts())
```

```
petal_length:
Null_values: 0
value counts:
```

|     |    |
|-----|----|
| 1.5 | 14 |
| 1.4 | 12 |
| 5.1 | 8  |
| 4.5 | 8  |
| 1.6 | 7  |
| 1.3 | 7  |
| 5.6 | 6  |
| 4.7 | 5  |
| 4.9 | 5  |
| 4.0 | 5  |
| 4.2 | 4  |
| 5.0 | 4  |
| 4.4 | 4  |
| 4.8 | 4  |
| 1.7 | 4  |
| 3.9 | 3  |
| 4.6 | 3  |
| 5.7 | 3  |
| 4.1 | 3  |
| 5.5 | 3  |
| 6.1 | 3  |
| 5.8 | 3  |
| 3.3 | 2  |
| 5.4 | 2  |
| 6.7 | 2  |
| 5.3 | 2  |
| 5.9 | 2  |
| 6.0 | 2  |
| 1.2 | 2  |
| 4.3 | 2  |
| 1.9 | 2  |
| 3.5 | 2  |
| 5.2 | 2  |
| 3.0 | 1  |
| 1.1 | 1  |
| 3.7 | 1  |
| 3.8 | 1  |
| 6.6 | 1  |
| 6.3 | 1  |
| 1.0 | 1  |
| 6.9 | 1  |
| 3.6 | 1  |
| 6.4 | 1  |

```
Name: petal_length, dtype: int64
```

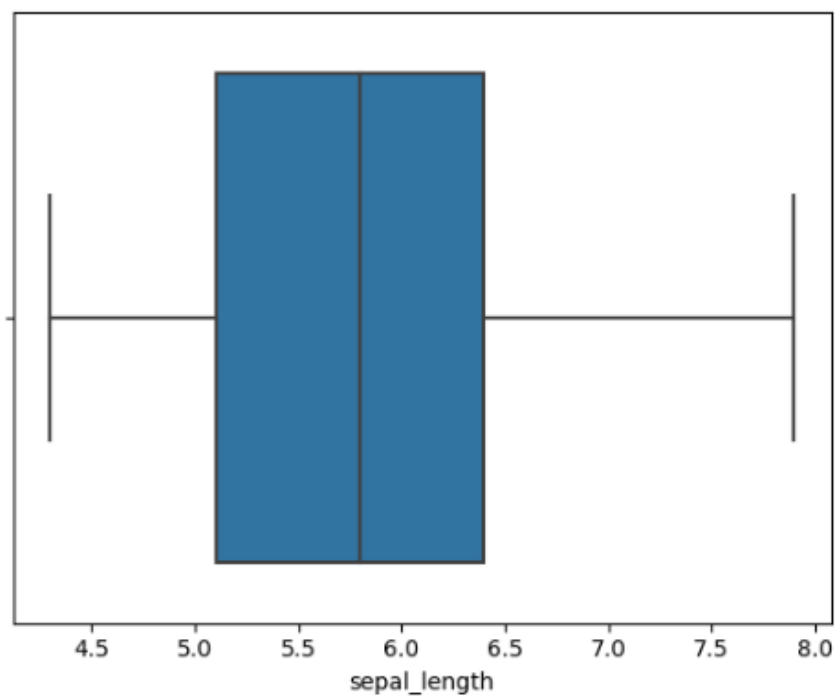
```
In [10]: print("petal_width:")
print("Null_values:",df['petal_width'].isna().sum())
print("value counts:\n",df['petal_width'].value_counts())
```

```
petal_width:
Null_values: 0
value counts:
 0.2      28
 1.3      13
 1.8      12
 1.5      12
 1.4       8
 2.3       8
 1.0       7
 0.4       7
 0.3       7
 0.1       6
 2.1       6
 2.0       6
 1.2       5
 1.9       5
 1.6       4
 2.5       3
 2.2       3
 2.4       3
 1.1       3
 1.7       2
 0.6       1
 0.5       1
Name: petal_width, dtype: int64
```

## OUTLIERS

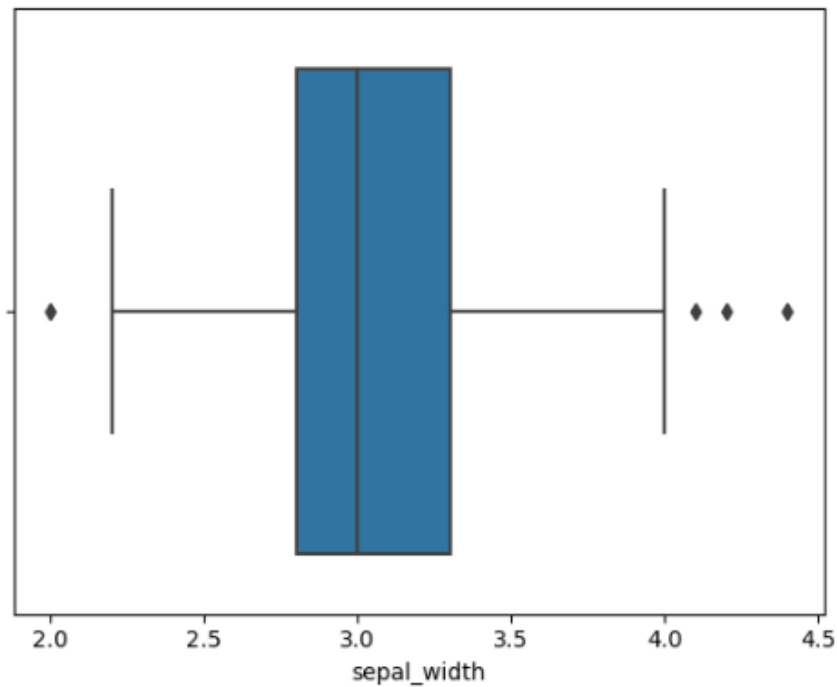
```
In [13]: sns.boxplot(df["sepal_length"])
plt.show
```

```
Out[13]: <function matplotlib.pyplot.show(close=None, block=None)>
```



```
In [14]: sns.boxplot(df["sepal_width"])
plt.show
```

```
Out[14]: <function matplotlib.pyplot.show(close=None, block=None)>
```



```
In [17]: q3 = df["sepal_width"].quantile(0.75)
q1 = df["sepal_width"].quantile(0.25)
iqr = q3-q1
iqr
```

```
Out[17]: 0.5
```

```
In [18]: upper_whisker = q3+1.5*iqr
lower_whisker = q1-1.5*iqr
```

```
In [19]: print(iqr)
print(upper_whisker)
print(lower_whisker)
```

```
0.5
4.05
2.05
```

```
In [20]: upper_whisker_value = df[(df["sepal_width"] > upper_whisker)].index
upper_whisker_value
```

```
Out[20]: Int64Index([15, 32, 33], dtype='int64')
```

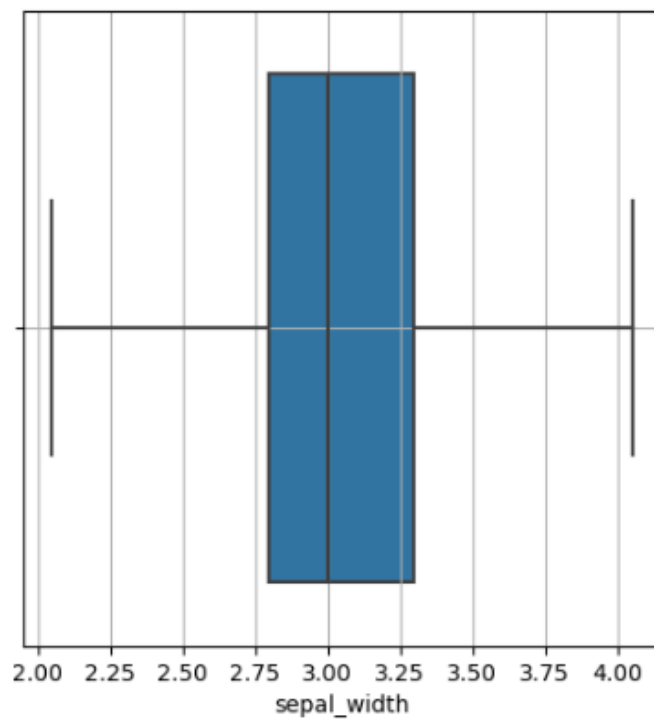
```
In [21]: lower_whisker_value = df[(df["sepal_width"] < lower_whisker)].index  
lower_whisker_value
```

```
Out[21]: Int64Index([60], dtype='int64')
```

```
In [22]: df.loc[upper_whisker_value,"sepal_width"] = np.nan  
df.fillna(upper_whisker,inplace=True)
```

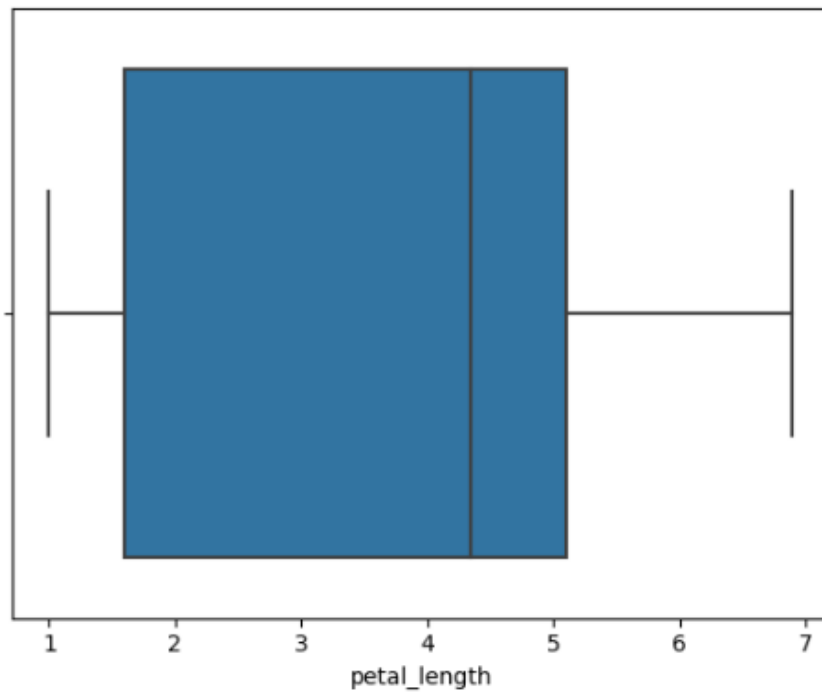
```
In [23]: df.loc[lower_whisker_value,"sepal_width"] = np.nan  
df.fillna(lower_whisker,inplace=True)
```

```
In [24]: plt.figure(figsize=(5,5))  
sns.boxplot(df['sepal_width'])  
plt.grid()
```



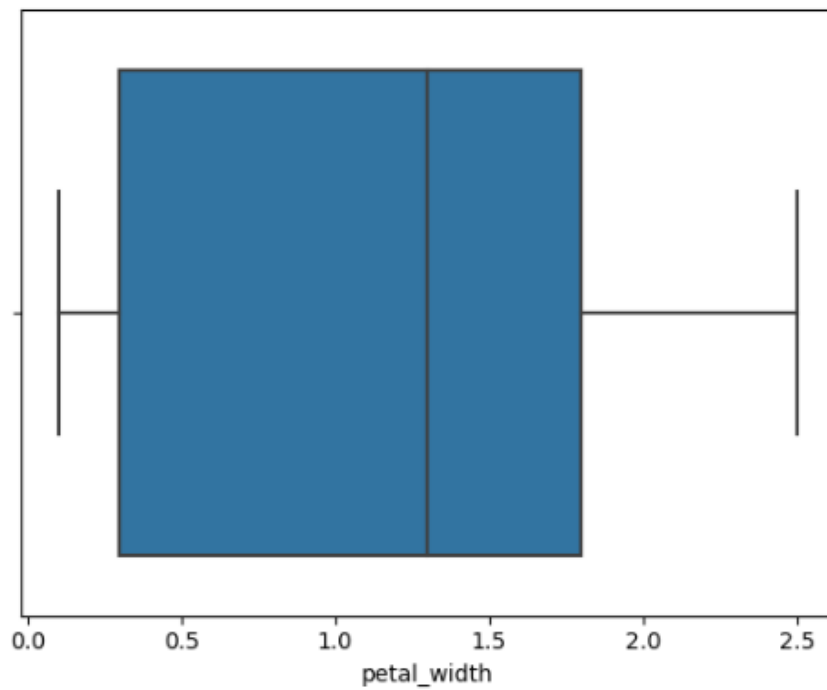
```
In [15]: sns.boxplot(df["petal_length"])  
plt.show
```

```
Out[15]: <function matplotlib.pyplot.show(close=None, block=None)>
```



```
In [16]: sns.boxplot(df["petal_width"])  
plt.show
```

```
Out[16]: <function matplotlib.pyplot.show(close=None, block=None)>
```

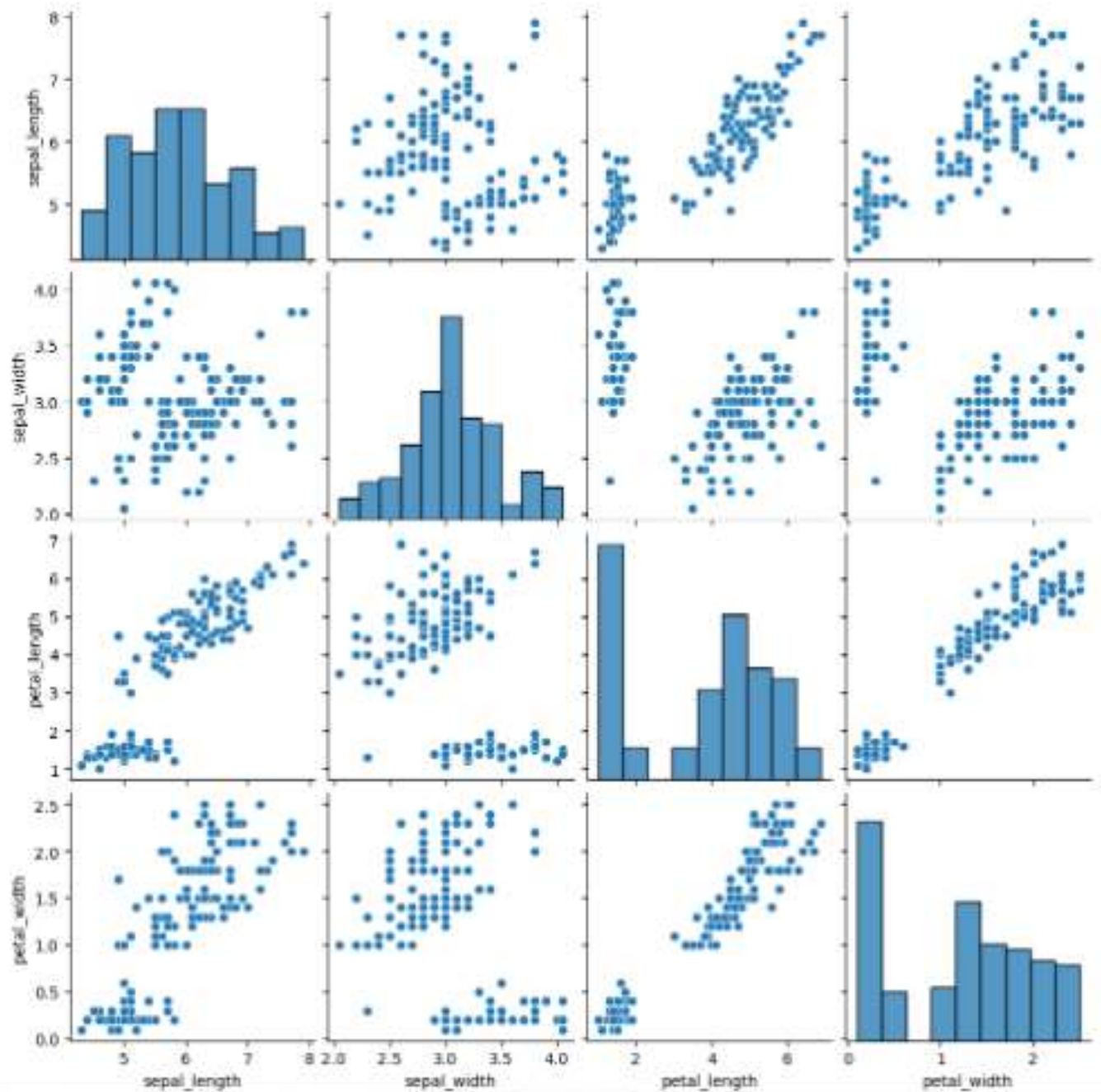




## PAIRPLOT

```
In [26]: sns.pairplot(df)
```

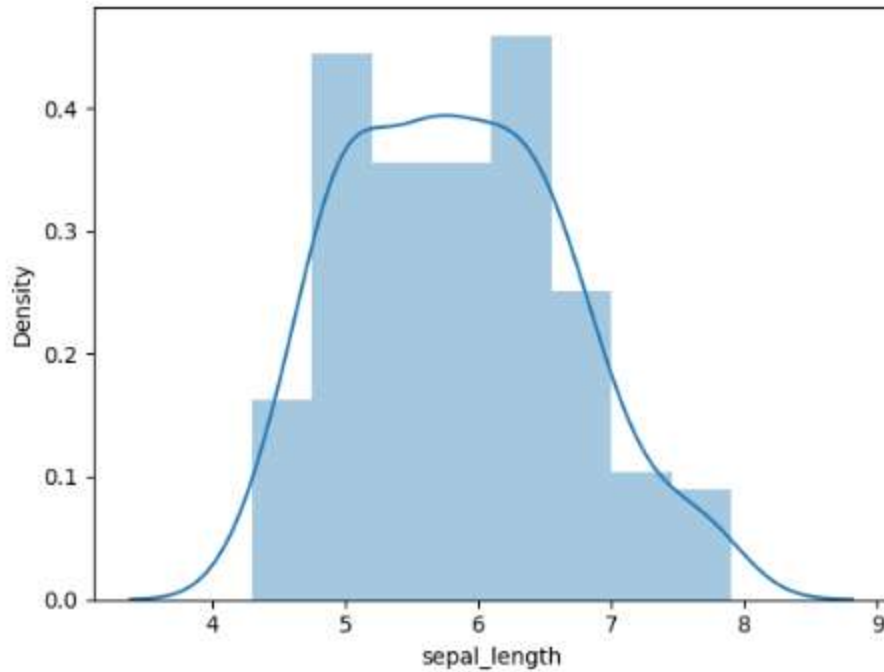
```
Out[26]: <matplotlib.axes._subplots.PairGrid at 0x1a123c78df8>
```



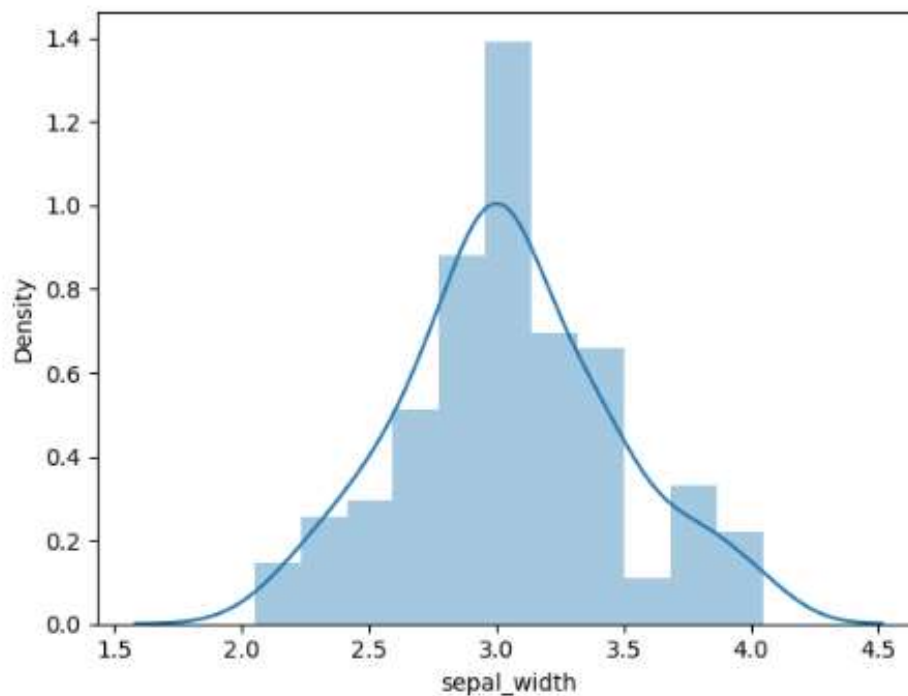
## SKEW

```
In [35]: from scipy.stats import skew
```

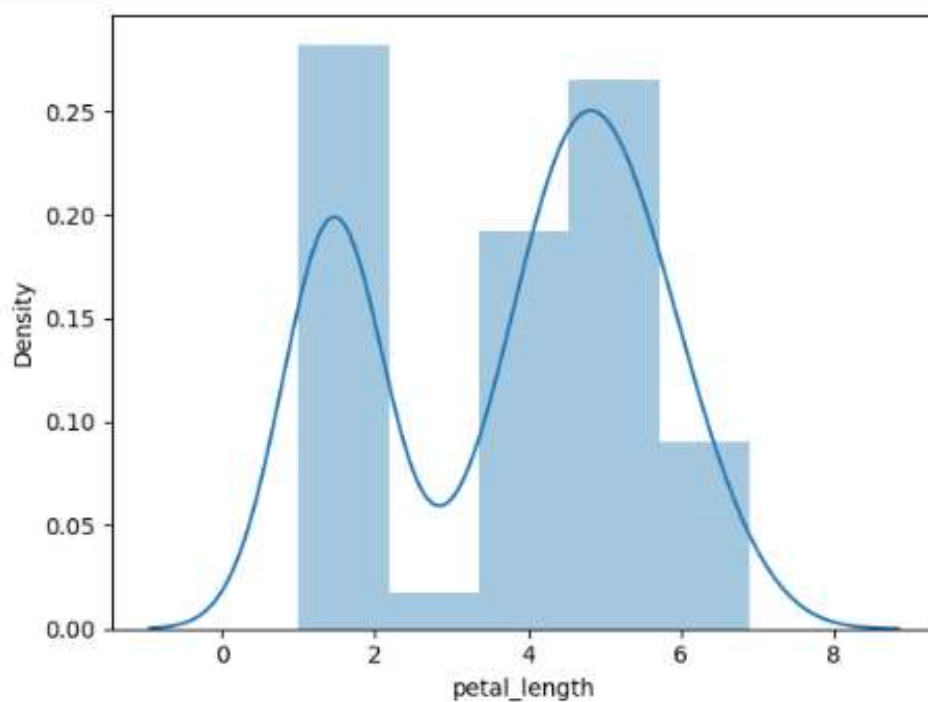
```
In [37]: skew(df['sepal_length'])  
sns.distplot(df['sepal_length'])  
plt.show()
```



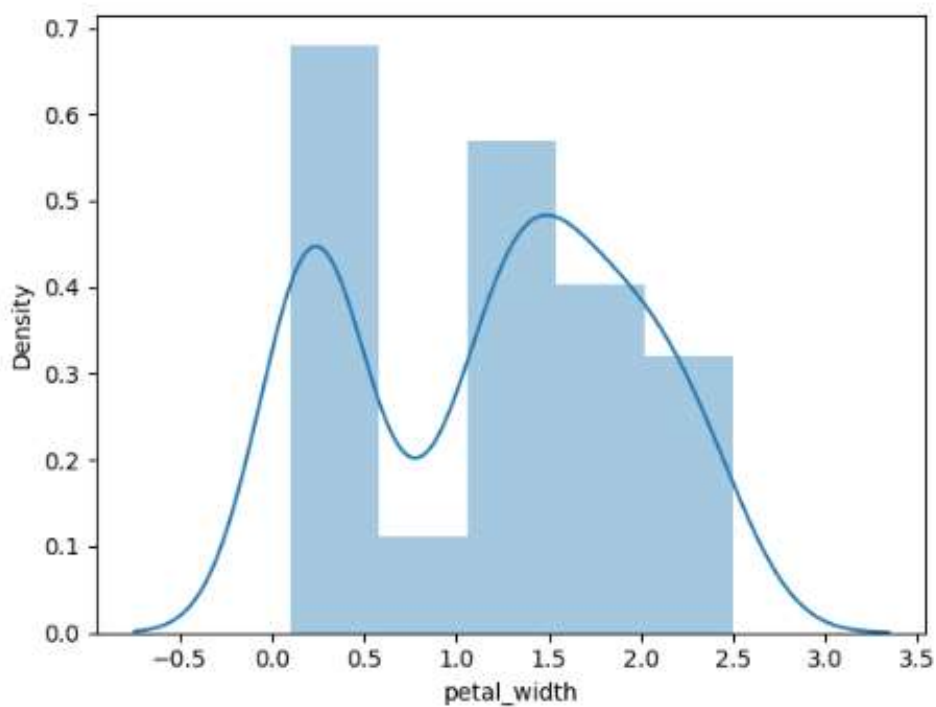
```
In [38]: skew(df['sepal_width'])  
sns.distplot(df['sepal_width'])  
plt.show()
```



```
In [39]: skew(df['petal_length'])
sns.distplot(df['petal_length'])
plt.show()
```



```
In [40]: skew(df['petal_width'])
sns.distplot(df['petal_width'])
plt.show()
```



## CORRELATION

```
In [50]: df.corr().style.background_gradient()
```

Out[50]:

|              | sepal_length | sepal_width | petal_length | petal_width | species   |
|--------------|--------------|-------------|--------------|-------------|-----------|
| sepal_length | 1.000000     | -0.110343   | 0.871754     | 0.817954    | 0.782561  |
| sepal_width  | -0.110343    | 1.000000    | -0.419823    | -0.355582   | -0.419264 |
| petal_length | 0.871754     | -0.419823   | 1.000000     | 0.962757    | 0.949043  |
| petal_width  | 0.817954     | -0.355582   | 0.962757     | 1.000000    | 0.956464  |
| species      | 0.782561     | -0.419264   | 0.949043     | 0.956464    | 1.000000  |

## ENCODING

```
In [42]: from sklearn.preprocessing import OrdinalEncoder
oe = OrdinalEncoder()
df['species'] = oe.fit_transform(df[['species']])
```

## SCALING

```
In [54]: x = df.select_dtypes("float64").columns
from sklearn.preprocessing import MinMaxScaler
sc = MinMaxScaler()
x = sc.fit_transform(x)
x = pd.DataFrame(x)
x.columns = ['sepal_length', 'sepal_width', 'petal_length', 'petal_width']
x
```

Out[54]:

|     | sepal_length | sepal_width | petal_length | petal_width |
|-----|--------------|-------------|--------------|-------------|
| 0   | 0.222222     | 0.725       | 0.067797     | 0.041667    |
| 1   | 0.166667     | 0.475       | 0.067797     | 0.041667    |
| 2   | 0.111111     | 0.575       | 0.050847     | 0.041667    |
| 3   | 0.083333     | 0.525       | 0.084746     | 0.041667    |
| 4   | 0.194444     | 0.775       | 0.067797     | 0.041667    |
| ... | ...          | ...         | ...          | ...         |
| 145 | 0.666667     | 0.475       | 0.711864     | 0.916667    |
| 146 | 0.555556     | 0.225       | 0.677966     | 0.750000    |
| 147 | 0.611111     | 0.475       | 0.711864     | 0.791667    |
| 148 | 0.527778     | 0.675       | 0.745763     | 0.916667    |
| 149 | 0.444444     | 0.475       | 0.694915     | 0.708333    |

150 rows × 4 columns

## FEATURE AND TARGET

```
In [55]: feature = x
target = df['species']
```

## MODEL BUILDING

```
In [56]: from sklearn.model_selection import train_test_split
xtrain,xtest,ytrain,ytest = train_test_split(feature,target,test_size=0.3,random_state=1)
```

```
In [57]: from sklearn.neighbors import KNeighborsClassifier
knn = KNeighborsClassifier(n_neighbors=5)
knn.fit(xtrain,ytrain)
```

```
Out[57]: KNeighborsClassifier()
```

```
In [58]: ypred = knn.predict(xtest)
ypred
```

```
Out[58]: array([0., 1., 1., 0., 2., 1., 2., 0., 0., 2., 1., 0., 2., 1., 1., 0., 1.,
        1., 0., 0., 1., 1., 2., 0., 2., 1., 0., 0., 1., 2., 1., 2., 1., 2.,
        2., 0., 1., 0., 1., 2., 2., 0., 1., 2., 1.])
```

```
In [62]: from sklearn.svm import SVC
svc = SVC()
svc.fit(xtrain,ytrain)
svcpred = svc.predict(xtest)
svcpred
```

```
Out[62]: array([0., 1., 1., 0., 2., 1., 2., 0., 0., 2., 1., 0., 2., 1., 1., 0., 1.,
        1., 0., 0., 1., 1., 2., 0., 2., 1., 0., 0., 1., 2., 1., 2., 1., 2.,
        2., 0., 1., 0., 1., 2., 2., 0., 1., 2., 1.])
```

## MODEL EVALUATION

```
In [60]: from sklearn.metrics import classification_report
```

```
In [66]: train = knn.score(xtrain, ytrain)
test = knn.score(xtest, ytest)
print("KNeighborsClassifier Report:")
print(f"Training Accuracy : {train}\nTesting Accuracy : {test}\n\n")
```

```
KNeighborsClassifier Report:
Training Accuracy : 0.9714285714285714
Testing Accuracy : 0.9555555555555556
```

```
In [67]: train = svc.score(xtrain, ytrain)
test = svc.score(xtest, ytest)
print("Support Vector Report:")
print(f"Training Accuracy : {train}\nTesting Accuracy : {test}\n\n")
```

```
Support Vector Report:
Training Accuracy : 0.9714285714285714
Testing Accuracy : 0.9555555555555556
```