```
In [1]: import numpy as np
          {\color{red}\textbf{import}} \  \, \text{pandas} \  \, {\color{red}\textbf{as}} \  \, \text{pd}
          import matplotlib.pyplot as plt
          import seaborn as sns
Out[10]:
                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
            145
                        6.7
                                    3.0
                                                5.2
                                                            2.3 virginica
           146
                        6.3
                                    2.5
                                                5.0
                                                            1.9 virginica
           147
                        6.5
                                                5.2
                                    3.0
                                                            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 [11]: data.head()
Out[11]:
              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
                                  3.1
                      4.6
                                              1.5
                                                          0.2
                                                               setosa
                      5.0
                                  3.6
                                              1.4
                                                          0.2
In [12]: data.describe()
Out[12]:
                  sepal_length sepal_width petal_length petal_width
                   150.000000
                               150.000000
                                           150.000000
                                                       150.000000
            mean
                     5.843333
                                 3.054000
                                             3.758667
                                                         1.198667
                     0.828066
                                 0.433594
                                             1.764420
                                                        0.763161
             std
             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 [13]: data.describe(include = 'object')
Out[13]:
                   species
            count
           unique
                        3
```

freq 50

In [14]: data.isnull().sum()

top

setosa

Out[14]: sepal_length 0 sepal_width 0 petal_length 0 petal_width 0 species 0 dtype: int64

```
In [15]: print("\n\nThe features in the dataset are as follows : ")
print("1. Sepal length : ", data['sepal_length'].dtype)
print("2. Sepal width : ", data['sepal_width'].dtype)
print("3. Petal length : ", data['petal_length'].dtype)
print("4. Petal width : ", data['petal_width'].dtype)
print("5. Species : ", data['species'].dtype)
```

```
The features in the dataset are as follows:

1. Sepal length: float64

2. Sepal width: float64

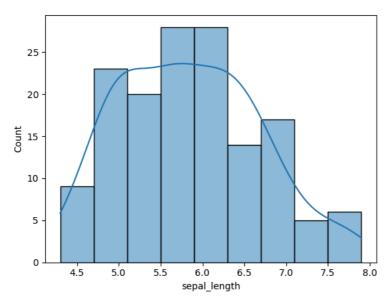
3. Petal length: float64

4. Petal width: float64

5. Species: object
```

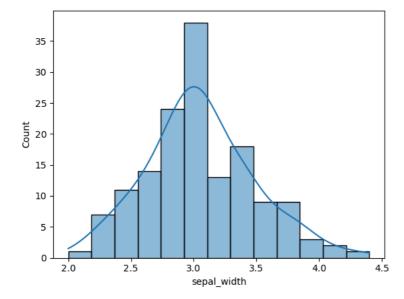
```
In [16]: sns.histplot(x = data['sepal_length'], kde=True)
```

Out[16]: <Axes: xlabel='sepal_length', ylabel='Count'>



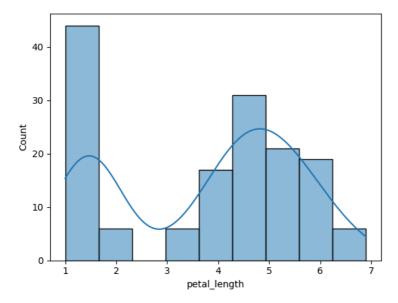
```
In [17]: sns.histplot(x = data['sepal_width'], kde=True)
```

Out[17]: <Axes: xlabel='sepal_width', ylabel='Count'>



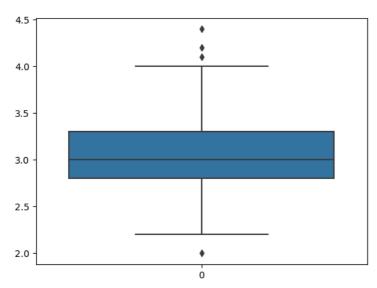
```
In [18]: sns.histplot(x = data['petal_length'], kde=True)
```

Out[18]: <Axes: xlabel='petal_length', ylabel='Count'>



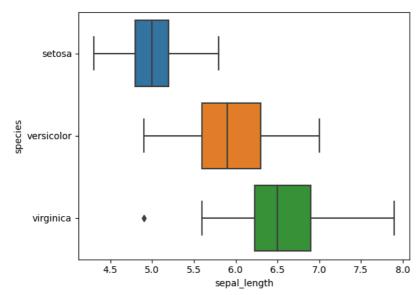
In [19]: sns.boxplot(data['sepal_width'])

Out[19]: <Axes: >



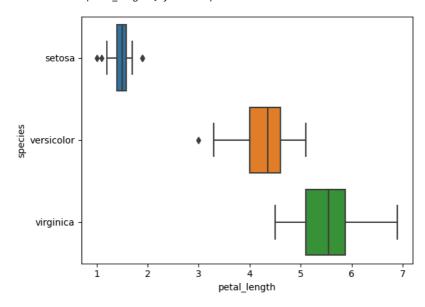
In [20]: sns.boxplot(x='sepal_length',y='species',data=data)

Out[20]: <Axes: xlabel='sepal_length', ylabel='species'>



```
In [21]: sns.boxplot(x='petal_length',y='species',data=data)
```

Out[21]: <Axes: xlabel='petal_length', ylabel='species'>



In []: