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# Assignment -10 -Data Visualization 3

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ERP Number: - 38

#### TE Comp 1

Download the Iris flower dataset or any other dataset into a DataFrame. (e.g., https://archive.ics.uci.edu/ml/datasets/Iris ). Scan the dataset and give the inference as:

- 1. List down the features and their types (e.g., numeric, nominal) available in the dataset.
- 2. Create a histogram for each feature in the dataset to illustrate the feature distributions.
- 3. Create a box plot for each feature in the dataset.
- 4. Compare distributions and identify outliers.

```
In [19]:
           import numpy as np
           import pandas as pd
           import matplotlib.pyplot as plt
           import seaborn as sns
In [20]:
           data=pd.read_csv('iris flower.csv')
In [21]:
           data.head()
Out[21]:
             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
                                                          0.2 Iris-setosa
                                  3.1
                                              1.5
                     5.0
                                  3.6
                                              1.4
                                                          0.2 Iris-setosa
In [22]:
           data.info()
          <class 'pandas.core.frame.DataFrame'>
          RangeIndex: 150 entries, 0 to 149
          Data columns (total 5 columns):
               Column
                              Non-Null Count Dtype
               sepal_length 150 non-null
                                                float64
           0
               sepal_width
                              150 non-null
                                                float64
           1
           2
               petal length 150 non-null
                                                float64
           3
               petal width
                              150 non-null
                                                float64
               species
                              150 non-null
                                                object
          dtypes: float64(4), object(1)
```

memory usage: 6.0+ KB

In [23]:

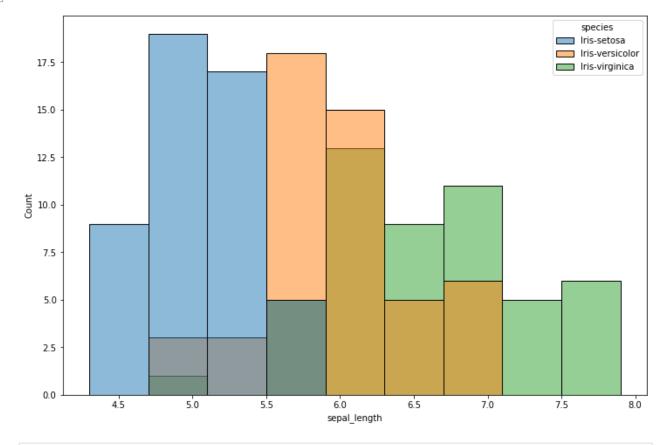
data.describe()

Οι	ıt	2	3	]	

	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

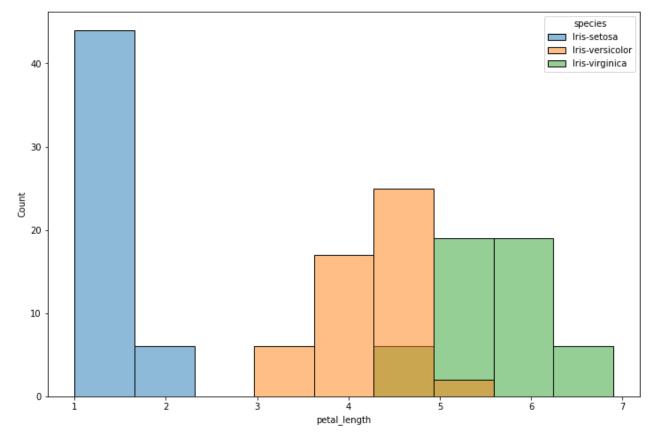
```
In [24]: plt.figure(figsize=(12,8))
    sns.histplot(x=data['sepal_length'], hue=data['species'])
    plt.plot()
```

Out[24]: []



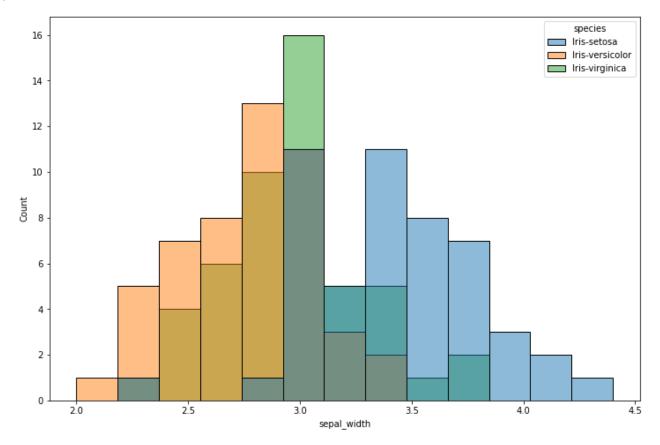
```
plt.figure(figsize=(12,8))
sns.histplot(x=data['petal_length'], hue=data['species'])
plt.plot()
```

Out[25]: []



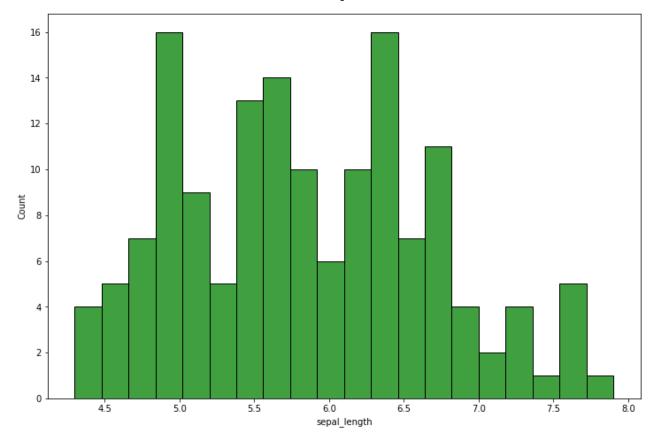
```
plt.figure(figsize=(12,8))
sns.histplot(x=data['sepal_width'], hue=data['species'])
plt.plot()
```

Out[26]: []



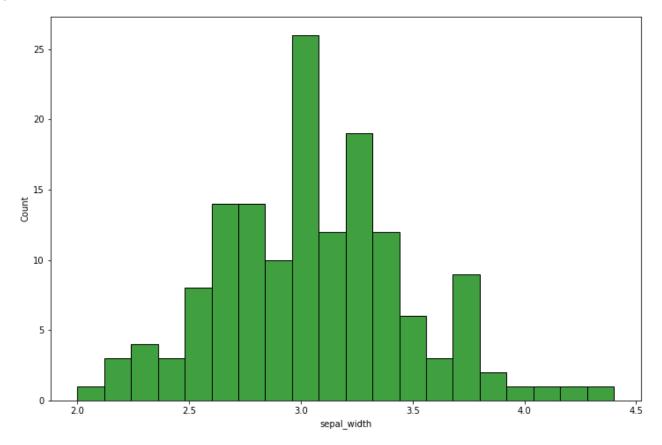
```
In [27]:
           plt.figure(figsize=(12,8))
            sns.histplot(x=data['petal_width'], hue=data['species'])
            plt.plot()
Out[27]: []
                                                                                                   species
                                                                                                 ■ Iris-setosa
             40
                                                                                                  Iris-versicolor
                                                                                                 ■ Iris-virginica
             35
             30
             25
             20
             15
             10
              5
             0.0
                                 0.5
                                                   1.0
                                                                     1.5
                                                                                       2.0
                                                           petal_width
In [28]:
            plt.figure(figsize=(12,8))
            sns.histplot(x=data['sepal_length'],bins=20, color='green')
            plt.plot()
```

Out[28]: []



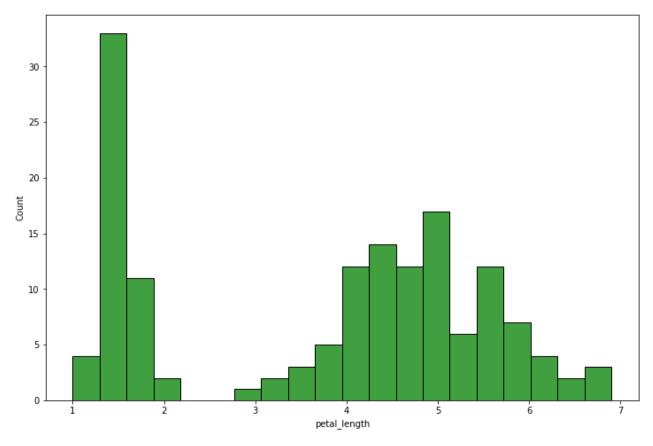
```
plt.figure(figsize=(12,8))
sns.histplot(x=data['sepal_width'],bins=20, color='green')
plt.plot()
```

Out[29]: []



```
In [30]: plt.figure(figsize=(12,8))
    sns.histplot(x=data['petal_length'],bins=20, color='green')
    plt.plot()
```

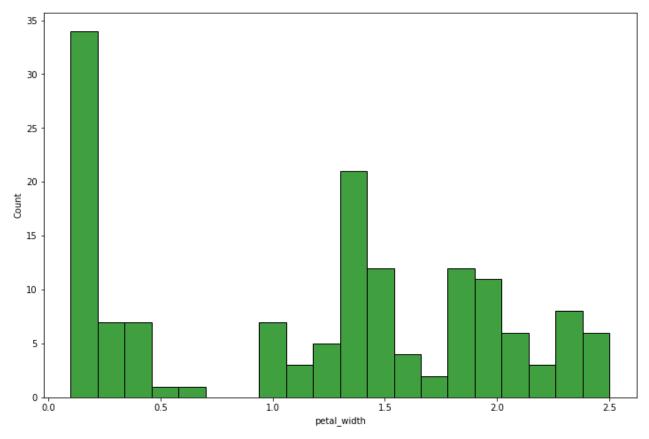
Out[30]: []



```
plt.figure(figsize=(12,8))
sns.histplot(x=data['petal_width'],bins=20, color='green')
plt.plot()
```

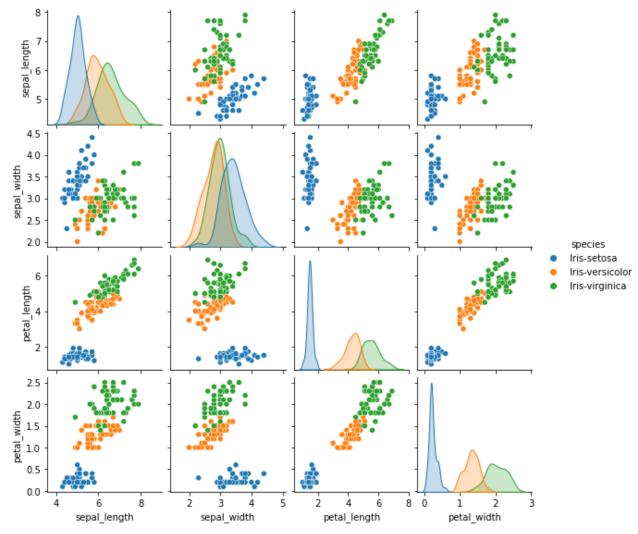
Out[31]: []

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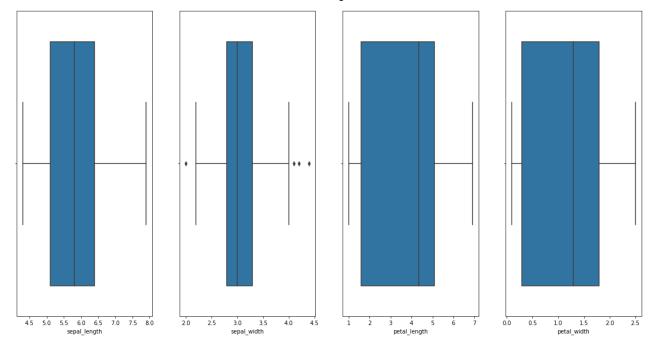


In [32]: sns.pairplot(data, hue='species', height=2)

Out[32]: <seaborn.axisgrid.PairGrid at 0x221433cf1c0>

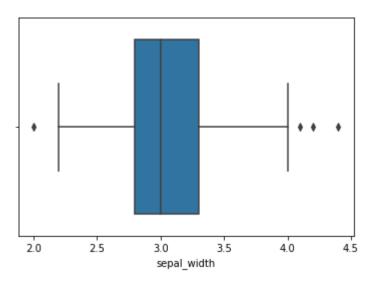


```
In [33]: import warnings
In [34]: warnings.filterwarnings('ignore')
    features_=data.columns.values[:-1]
    fig=plt.figure(figsize=(20,10))
    for columns, feature in enumerate(features_):
        fig.add_subplot(1,4,columns+1)
        sns.boxplot(data[feature],data=data)
    plt.show()
```



```
In [38]: sns.boxplot(x='sepal_width', data=data)
```

### Out[38]: <AxesSubplot:xlabel='sepal\_width'>



In the above graph, the values above 4 and below 2 are acting as outliers.

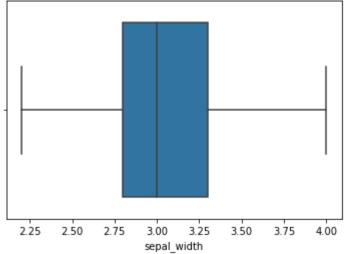
```
In [44]:
# Removing Outliers
# IQR
Q1 = np.percentile(data['sepal_width'], 25,interpolation = 'midpoint')
Q3 = np.percentile(data['sepal_width'], 75,interpolation = 'midpoint')
IQR = Q3 - Q1
print("Old Shape: ", data.shape)
# Upper bound
upper = np.where(data['sepal_width'] >= (Q3+1.5*IQR))
# Lower bound
lower = np.where(data['sepal_width'] <= (Q1-1.5*IQR))</pre>
```

```
# Removing the Outliers
data.drop(upper[0], inplace = True)
data.drop(lower[0], inplace = True)
print("New Shape: ", data.shape)
sns.boxplot(x='sepal_width', data=data)
```

Old Shape: (150, 5) New Shape: (146, 5)

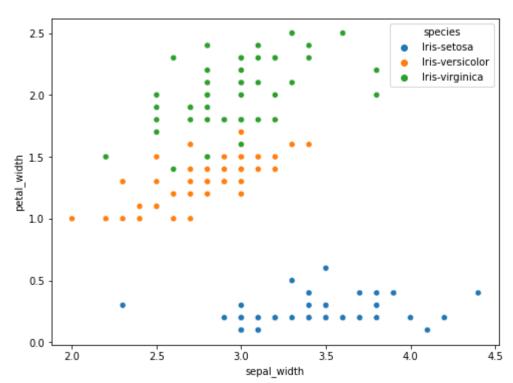
Out[44]: <AxesSubplot:xlabel='sepal\_width'>

## Out[44].



```
In [35]:
    plt.figure(figsize=(8,6))
    sns.scatterplot(data=data,x='sepal_width', y='petal_width', hue='species')
    plt.plot()
```

## Out[35]: []



```
In [36]:
    plt.figure(figsize=(8,6))
    sns.scatterplot(data=data,x='sepal_length', y='petal_length', hue='species')
    plt.plot()
```

Out[36]: []

