

In []:

```
Task 1: PERFORM DATA CLEANING
clean a dataset by removing missing values and outliers
By NIVEDHA M
```

In []:

```
#IMPLEMENTING THE DEPENDENCIES
```

In [1]:

```
import pandas as pd
import numpy as np
iTask 1: PERFORM DATA CLEANING
clean a dataset by removing missing values and outliers
By NIVEDHA Mimport seaborn as sns
```

In [12]:

```
#DATA READING
```

In [2]:

```
gender_data = pd.read_csv("gender_submission.csv")
print(gender_data)
```

| | PassengerId | Survived |
|-----|-------------|----------|
| 0 | 892 | 0 |
| 1 | 893 | 1 |
| 2 | 894 | 0 |
| 3 | 895 | 0 |
| 4 | 896 | 1 |
| .. | ... | ... |
| 413 | 1305 | 0 |
| 414 | 1306 | 1 |
| 415 | 1307 | 0 |
| 416 | 1308 | 0 |
| 417 | 1309 | 0 |

[418 rows x 2 columns]

In []:

```
#DATA CLEANING
#Fill the missing values for passenger id and survival columns.In order to fill the miss
#will fill the missing values of both the columns by taking the mean of all columns
```

In [8]:

```
#fill passengerID column
gender_data["PassengerId"].fillna(gender_data["PassengerId"].mean(),inplace = True)
gender_data["PassengerId"].isna().sum()
```

Out[8]:

0

In [9]:

```
#fill survived column
gender_data["Survived"].fillna(gender_data["Survived"].mean(), inplace=True)
gender_data["Survived"].isna().sum()
```

Out[9]:

0

In []:

```
#Alternatively we will visualize the null value using heatmap
#we will use heatmap method by passing only records which are null
```

In [10]:

```
sns.heatmap(gender_data.isna())
```

Out[10]:

<AxesSubplot:>



In []:

```
#we can conclude from the above heatmap that there is no null value left in our dataset
```

In []:

Task 2 : Calculate Summary Statistics

Calculate summary statistics(mean, median, mode, standard deviation) for a dataset
By NIVEDHA M

In []:

#Implementing the Dependencies

In [1]:

```
import pandas as pd
import numpy as np
```

In []:

#Data Reading

In [2]:

```
gender_data = pd.read_csv("gender_submission.csv")
print(gender_data)
```

| | PassengerId | Survived |
|-----|-------------|----------|
| 0 | 892 | 0 |
| 1 | 893 | 1 |
| 2 | 894 | 0 |
| 3 | 895 | 0 |
| 4 | 896 | 1 |
| .. | ... | ... |
| 413 | 1305 | 0 |
| 414 | 1306 | 1 |
| 415 | 1307 | 0 |
| 416 | 1308 | 0 |
| 417 | 1309 | 0 |

[418 rows x 2 columns]

In []:

#Using the describe() to find the statistics(mean, median, mode, standard deviation)

In [3]:

```
#Calculating the statistics (mean, median, mode, standard deviation)
gender_data.describe()
```

Out[3]:

| | PassengerId | Survived |
|-------|-------------|------------|
| count | 418.000000 | 418.000000 |
| mean | 1100.500000 | 0.363636 |
| std | 120.810458 | 0.481622 |
| min | 892.000000 | 0.000000 |
| 25% | 996.250000 | 0.000000 |
| 50% | 1100.500000 | 0.000000 |
| 75% | 1204.750000 | 1.000000 |
| max | 1309.000000 | 1.000000 |

In [5]:

```
gender_data.median()
```

Out[5]:

PassengerId 1100.5
Survived 0.0
dtype: float64

In [6]:

```
gender_data.mode()
```

Out[6]:

| | PassengerId | Survived |
|-----|-------------|----------|
| 0 | 892 | 0.0 |
| 1 | 893 | NaN |
| 2 | 894 | NaN |
| 3 | 895 | NaN |
| 4 | 896 | NaN |
| ... | ... | ... |
| 413 | 1305 | NaN |
| 414 | 1306 | NaN |
| 415 | 1307 | NaN |
| 416 | 1308 | NaN |
| 417 | 1309 | NaN |

418 rows × 2 columns

In []:

```
#TASK 3 : Visualization using Histogram
```

```
#Create a histogram or bar chart to visualize the distribution of data in a dataclasses_to_dicts
```

```
#By NIVEDHA M
```

```
#Implementing the Dependencies
```

```
import pandas as pd
import seaborn as sns
```

```
#Reading the datasets
```

```
iris_data = pd.read_csv("Iris.csv")
print(iris_data)
```

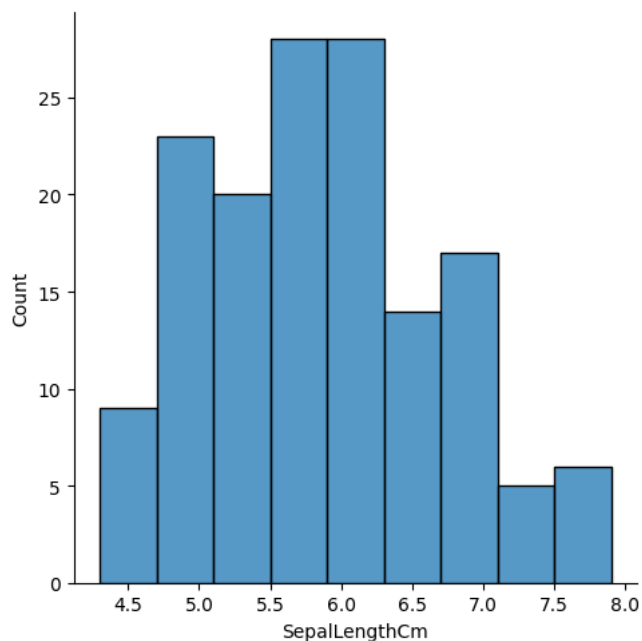
| | Id | SepalLengthCm | SepalWidthCm | PetalLengthCm | PetalWidthCm | \ |
|-----|-----|---------------|--------------|---------------|--------------|---|
| 0 | 1 | 5.1 | 3.5 | 1.4 | 0.2 | |
| 1 | 2 | 4.9 | 3.0 | 1.4 | 0.2 | |
| 2 | 3 | 4.7 | 3.2 | 1.3 | 0.2 | |
| 3 | 4 | 4.6 | 3.1 | 1.5 | 0.2 | |
| 4 | 5 | 5.0 | 3.6 | 1.4 | 0.2 | |
| .. | ... | ... | ... | ... | ... | |
| 145 | 146 | 6.7 | 3.0 | 5.2 | 2.3 | |
| 146 | 147 | 6.3 | 2.5 | 5.0 | 1.9 | |
| 147 | 148 | 6.5 | 3.0 | 5.2 | 2.0 | |
| 148 | 149 | 6.2 | 3.4 | 5.4 | 2.3 | |
| 149 | 150 | 5.9 | 3.0 | 5.1 | 1.8 | |

| | Species |
|-----|----------------|
| 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 |

```
[150 rows x 6 columns]
```

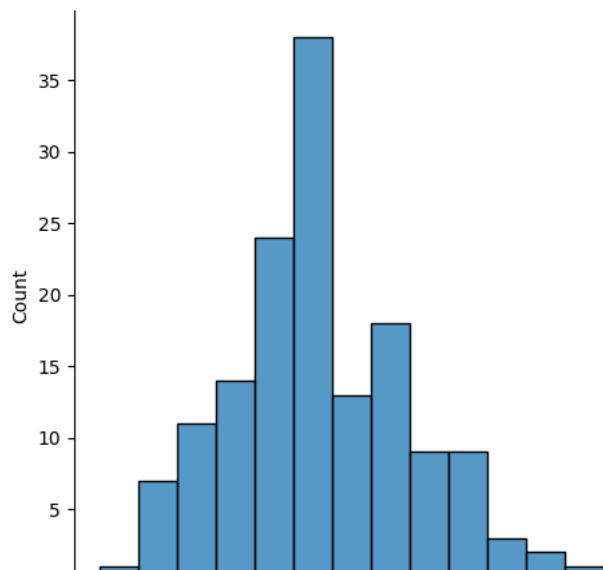
```
#plotting the histogram for SepalLength
sns.displot(x = "SepalLengthCm", data = iris_data)
```

```
<seaborn.axisgrid.FacetGrid at 0x78389ae7aef0>
```



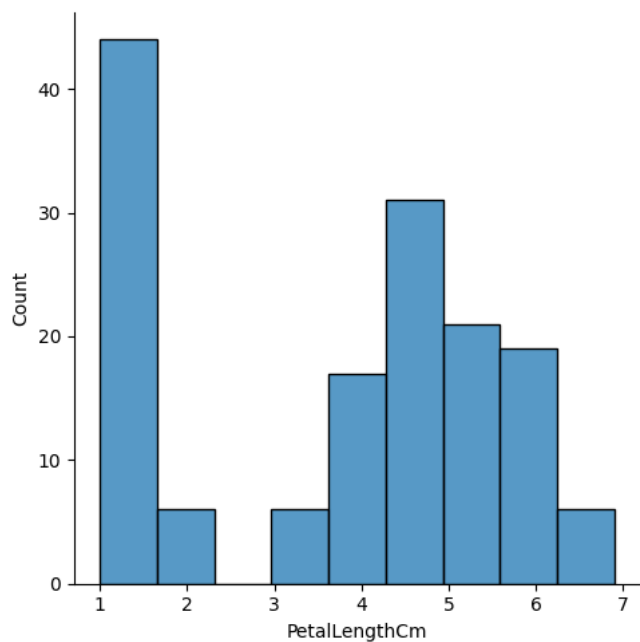
```
#plotting the histogram for SepalWidth
sns.displot(x = "SepalWidthCm", data = iris_data)
```

```
<seaborn.axisgrid.FacetGrid at 0x7838d2440d30>
```



```
#plotting the histogram for PetalLength  
sns.displot(x = "PetalLengthCm", data = iris_data)
```

```
<seaborn.axisgrid.FacetGrid at 0x78389a8c7ac0>
```



```
#plotting the histogram for PetalWidth  
sns.displot(x = "PetalWidthCm", data = iris_data)
```

