```
import numpy as np
import matplotlib.pyplot as plt
import pandas as pd
import seaborn as sns

df = pd.read_csv('iris.csv')
df.head()
```

	Id	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm	Species
0	1	5.1	3.5	1.4	0.2	setosa
1	2	4.9	3.0	1.4	0.2	setosa
2	3	4.7	3.2	1.3	0.2	setosa
3	4	4.6	3.1	1.5	0.2	setosa
4	5	5.0	3.6	1.4	0.2	setosa

How many features are there and what are their types (e.g., numeric, nominal)?

```
df.info()
RangeIndex: 150 entries, 0 to 149
     Data columns (total 6 columns):
     # Column
                    Non-Null Count Dtype
     0 Id
                       150 non-null
                                       int64
     1 SepalLengthCm 150 non-null float64
     2 SepalWidthCm 150 non-null float64
3 PetalLengthCm 150 non-null float64
     4 PetalWidthCm 150 non-null float64
5 Species 150 non-null object
     dtypes: float64(4), int64(1), object(1)
     memory usage: 7.2+ KB
np.unique(df["Species"])
     array(['setosa', 'versicolor', 'virginica'], dtype=object)
```

Compute and display summary statistics for each feature available in the dataset.(eg.

minimum value, maximum value, mean, range, standard deviation, variance and percentiles

df.describe()

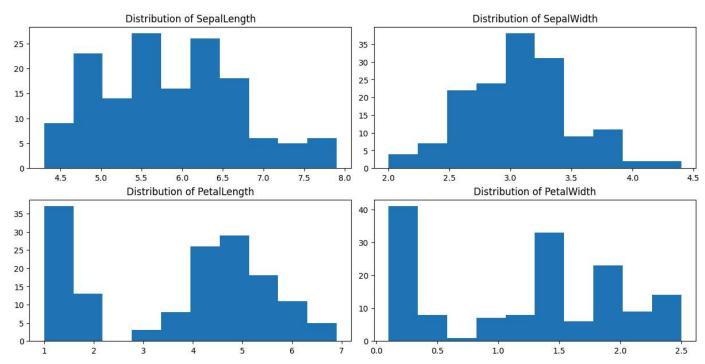
	Id	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm
count	150.000000	150.000000	150.000000	150.000000	150.000000
mean	75.500000	5.843333	3.054000	3.758667	1.198667
std	43.445368	0.828066	0.433594	1.764420	0.763161
min	1.000000	4.300000	2.000000	1.000000	0.100000
25%	38.250000	5.100000	2.800000	1.600000	0.300000
50%	75.500000	5.800000	3.000000	4.350000	1.300000
75%	112.750000	6.400000	3.300000	5.100000	1.800000
max	150.000000	7.900000	4.400000	6.900000	2.500000

Data Visualization-Create a histogram for each feature in the dataset to illustrate the feature distributions

```
import seaborn as sns
import matplotlib
import matplotlib.pyplot as plt

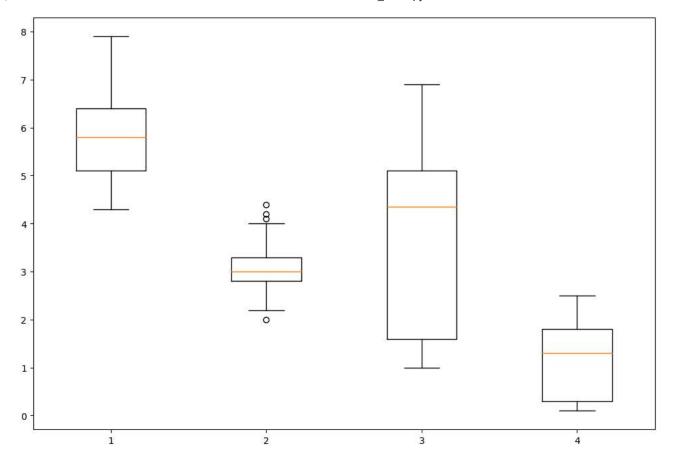
fig, axes = plt.subplots(2, 2, figsize=(12, 6), constrained_layout = True)

for i in range(4):
    x, y = i // 2, i % 2
    _ = axes[x, y].hist(df[df.columns[i + 1]])
    _ = axes[x, y].set_title(f"Distribution of {df.columns[i + 1][:-2]}")
```



Create a boxplot for each feature in the dataset. All of the boxplots should be combined into a single plot. Compare distributions and identify outliers.

```
data_to_plot = df[df.columns[1:-1]]
fig, axes = plt.subplots(1, figsize=(12,8))
bp = axes.boxplot(data_to_plot)
```



If we observe closely for the box 2, interquartile distance is roughly around 0.75 hence the values lying beyond this range of (third quartile + interquartile distance) i.e. roughly around 4.05 will be considered as outliers. Similarly outliers with other boxplots can be found.