

Assignment 3

1. Take any dataset of your own from Kaggle/Kdnuggets, apply Data analytics and Data visualization using Pandas, Matplotlib and Seaborn . Do various kinds of EDA analytics.



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[3]:

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

# Load the dataset
df = pd.read_csv('/kaggle/input/iris-flower-dataset/IRIS.csv')

# Display the first few rows of the dataframe
print("First few rows of the dataset:")
print(df.head())
```

First few rows of the dataset:

	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

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[4]:

```
# Summary statistics
print("\nSummary statistics:")
print(df.describe())
```

Summary statistics:

	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





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[4]:

```
# Summary statistics
print("\nSummary statistics:")
print(df.describe())
```

```
Summary statistics:
   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
```

[5]:

```
# Check for missing values
print("\nMissing values:")
print(df.isnull().sum())
```

```
Missing values:
sepal_length    0
sepal_width     0
petal_length    0
petal_width     0
species         0
dtype: int64
```

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[6]:

```
# Distribution of the target variable
```



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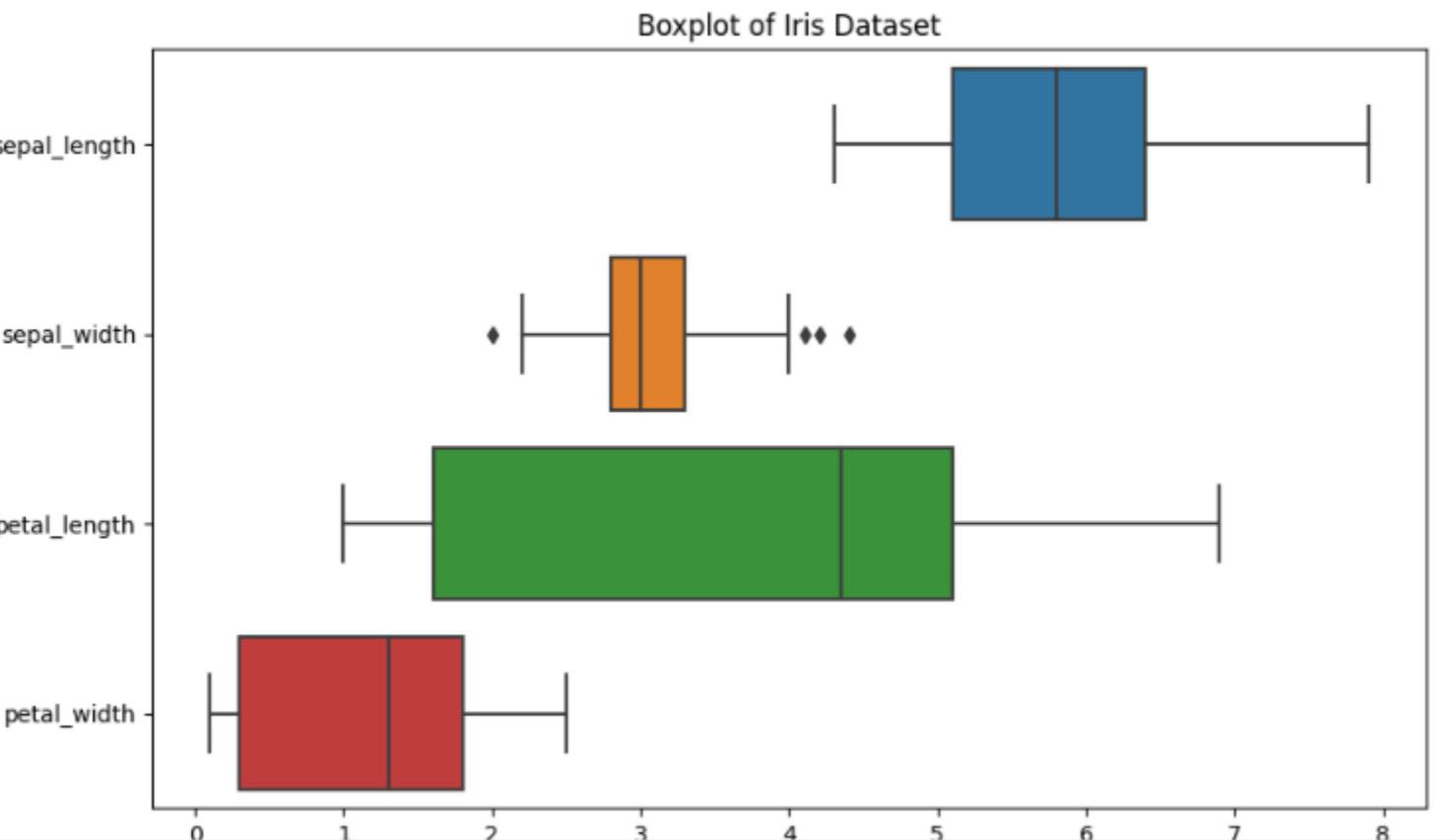
[6]:

```
# Distribution of the target variable
print("\nDistribution of the target variable:")
print(df['species'].value_counts())
```

```
Distribution of the target variable:
species
Iris-setosa      50
Iris-versicolor  50
Iris-virginica   50
Name: count, dtype: int64
```

[11]:

```
# Boxplot
plt.figure(figsize=(10, 6))
sns.boxplot(data=df, orient='h')
plt.title('Boxplot of Iris Dataset')
plt.show()
```





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Code

```
# Histograms
plt.figure(figsize=(10, 6))
for i, feature in enumerate(df.columns[:-1]):
    plt.subplot(2, 2, i+1)
    sns.histplot(data=df, x=feature, kde=True)
    plt.title(f'Histogram of {feature}')
plt.tight_layout()
plt.show()
```

```
/opt/conda/lib/python3.10/site-packages/seaborn/_oldcore.py:1119: FutureWarning: use_inf_as_na option is deprecated and will be removed in a future version. Convert inf values to NaN before operating instead.
  with pd.option_context('mode.use_inf_as_na', True):
/opt/conda/lib/python3.10/site-packages/seaborn/_oldcore.py:1119: FutureWarning: use_inf_as_na option is deprecated and will be removed in a future version. Convert inf values to NaN before operating instead.
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```

