

practical-10-piyusha-supe

April 12, 2025

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Data Visualization III

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 boxplot for each feature in the dataset. 4. Compare distributions and identify outliers

```
[1]: # Required Libraries
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt

# Load the Iris Dataset (from seaborn)
iris = sns.load_dataset("iris")

# Preview the dataset
print("Dataset:")
print(iris.head())
print(iris.tail())
print(iris.info())
print(iris.describe(include = "all"))
print(iris.shape)
print(iris.size)
print(iris.ndim)
print(iris.columns)
print(iris.dtypes)
```

Dataset:

| | 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 | setosa |

| | sepal_length | sepal_width | petal_length | petal_width | species |
|-----|--------------|-------------|--------------|-------------|-----------|
| 145 | 6.7 | 3.0 | 5.2 | 2.3 | virginica |

```

146      6.3      2.5      5.0      1.9  virginica
147      6.5      3.0      5.2      2.0  virginica
148      6.2      3.4      5.4      2.3  virginica
149      5.9      3.0      5.1      1.8  virginica
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 150 entries, 0 to 149
Data columns (total 5 columns):
 #   Column        Non-Null Count  Dtype  
--- 
 0   sepal_length  150 non-null    float64
 1   sepal_width   150 non-null    float64
 2   petal_length  150 non-null    float64
 3   petal_width   150 non-null    float64
 4   species       150 non-null    object  
dtypes: float64(4), object(1)
memory usage: 6.0+ KB
None
      sepal_length  sepal_width  petal_length  petal_width  species
count    150.000000  150.000000  150.000000  150.000000  150
unique     NaN         NaN         NaN         NaN         3
top        NaN         NaN         NaN         NaN         setosa
freq       NaN         NaN         NaN         NaN         50
mean      5.843333  3.057333  3.758000  1.199333  NaN
std       0.828066  0.435866  1.765298  0.762238  NaN
min       4.300000  2.000000  1.000000  0.100000  NaN
25%      5.100000  2.800000  1.600000  0.300000  NaN
50%      5.800000  3.000000  4.350000  1.300000  NaN
75%      6.400000  3.300000  5.100000  1.800000  NaN
max      7.900000  4.400000  6.900000  2.500000  NaN
(150, 5)
750
2
Index(['sepal_length', 'sepal_width', 'petal_length', 'petal_width',
       'species'],
      dtype='object')
sepal_length    float64
sepal_width     float64
petal_length    float64
petal_width     float64
species        object
dtype: object

```

```
[2]: # 1. Feature List and Types
print("\nFeature Types:")
for column in iris.columns:
    dtype = iris[column].dtype
    ftype = "Nominal" if dtype == "object" else "Numeric"
```

```

print(f"{column}: {ftype} ({dtype})")

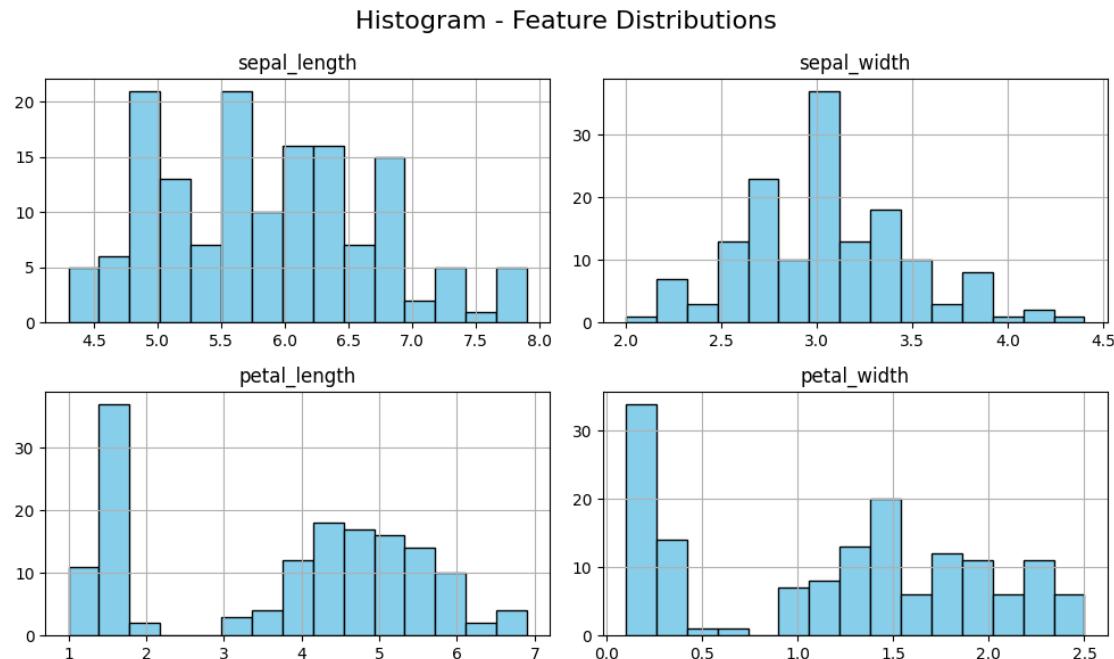
# 2. Histograms for Each Numeric Feature
iris_numeric = iris.drop(columns='species') # Only numeric features

iris_numeric.hist(bins=15, figsize=(10, 6), color='skyblue', edgecolor='black')
plt.suptitle("Histogram - Feature Distributions", fontsize=16)
plt.tight_layout()
plt.show()

```

Feature Types:

- sepal_length: Numeric (float64)
- sepal_width: Numeric (float64)
- petal_length: Numeric (float64)
- petal_width: Numeric (float64)
- species: Nominal (object)



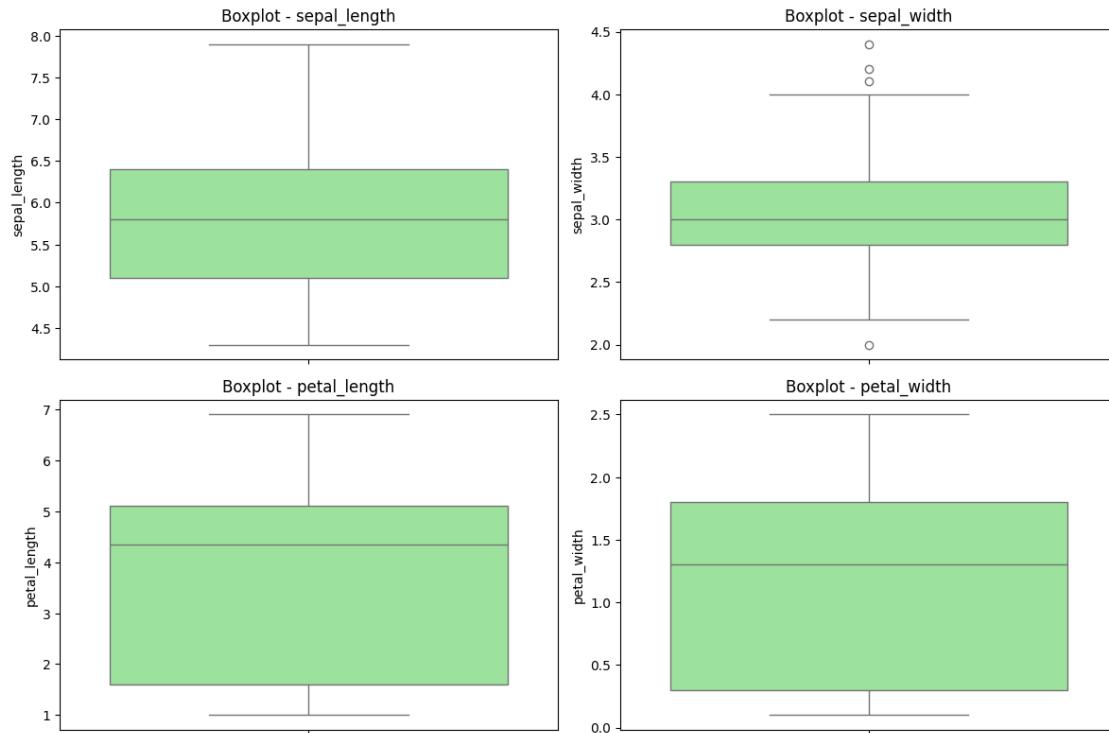
[3]: # 3. Boxplots for Each Feature

```

plt.figure(figsize=(12, 8))
for i, col in enumerate(iris_numeric.columns, 1):
    plt.subplot(2, 2, i)
    sns.boxplot(y=iris[col], color='lightgreen')
    plt.title(f"Boxplot - {col}")
plt.tight_layout()

```

```
plt.show()
```



```
[5]: # 4. Observations
print("\nObservations & Inference:")

# Describe statistics
print("\nDescriptive Statistics:")
print(iris_numeric.describe())

# Check for outliers using IQR method
print("\nOutliers (using IQR):")
for column in iris_numeric.columns:
    Q1 = iris[column].quantile(0.25)
    Q3 = iris[column].quantile(0.75)
    IQR = Q3 - Q1
    lower = Q1 - 1.5 * IQR
    upper = Q3 + 1.5 * IQR
    outliers = iris[(iris[column] < lower) | (iris[column] > upper)]
    print(f"{column}: {len(outliers)} outlier(s)")
```

Observations & Inference:

Descriptive Statistics:

| | sepal_length | sepal_width | petal_length | petal_width |
|-------|--------------|-------------|--------------|-------------|
| count | 150.000000 | 150.000000 | 150.000000 | 150.000000 |
| mean | 5.843333 | 3.057333 | 3.758000 | 1.199333 |
| std | 0.828066 | 0.435866 | 1.765298 | 0.762238 |
| 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 |

Outliers (using IQR):

sepal_length: 0 outlier(s)
sepal_width: 4 outlier(s)
petal_length: 0 outlier(s)
petal_width: 0 outlier(s)

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