

## Code Explanation

### 1. Importing Libraries

The code begins by importing the necessary libraries:

python

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```
import numpy as np

import pandas as pd

import matplotlib.pyplot as plt

%matplotlib inline

import seaborn as sns
```

- **numpy**: A library for numerical computations.
- **pandas**: Used for data manipulation and analysis, particularly to handle DataFrames.
- **matplotlib.pyplot**: A library for plotting graphs and visualizations.
- **%matplotlib inline**: This is a magic command to display the plots inline in Jupyter notebooks.
- **seaborn**: A data visualization library based on matplotlib that provides a higher-level interface for creating informative statistical plots.

### 2. Loading the Iris Dataset

The dataset is loaded using the seaborn library:

python

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```
dataset = sns.load_dataset('iris')
```

- The `sns.load_dataset('iris')` function loads the famous Iris flower dataset, which contains information about the lengths and widths of the sepals and petals for three different species of Iris flowers.

### 3. Exploring the Dataset

The following code snippet shows a preview of the dataset and its data types:

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```
dataset.head()
```

```
dataset.info()
```

- `dataset.head()` displays the first few rows of the dataset, allowing you to get an overview of its structure.
- `dataset.info()` provides information about the DataFrame, such as the number of non-null entries, column data types, and memory usage.

### 4. Histograms for Each Feature

The next block of code creates histograms for the features of the dataset:

python

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```
fig, axes = plt.subplots(2, 2, figsize=(16, 8))
```

```
axes[0,0].set_title("Distribution of Sepal Length")
```

```
axes[0,0].hist(dataset["sepal_length"]);
```

```
axes[0,1].set_title("Distribution of Sepal Width")
```

```
axes[0,1].hist(dataset["sepal_width"]);
```

```
axes[1,0].set_title("Distribution of Petal Length")
```

```
axes[1,0].hist(dataset["petal_length"]);
```

```
axes[1,1].set_title("Distribution of Petal Width")
```

```
axes[1,1].hist(dataset["petal_width"]);
```

- `fig, axes = plt.subplots(2, 2, figsize=(16, 8))`: This line creates a 2x2 grid of subplots (four total), with a specified figure size of (16, 8).
- The next lines plot histograms for each of the four features in the dataset (`sepal_length`, `sepal_width`, `petal_length`, and `petal_width`) in separate subplots.
- `axes[x, y].set_title()` is used to set the title for each subplot, while `axes[x, y].hist()` plots the histogram for the respective feature.

## 5. Boxplots for Each Feature

The next section of the code creates boxplots for each feature with respect to the species of the Iris flower:

```
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```

```
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```

```
fig, axes = plt.subplots(2, 2, figsize=(16,9))
```

```
axes[0,0].set_title("Distribution of Sepal Length")
```

```
sns.boxplot(y="sepal_length", x="species", data=dataset,  
orient='v', ax=axes[0, 0])
```

```
axes[0,1].set_title("Distribution of Sepal Width")

sns.boxplot(y="sepal_width", x="species", data=dataset,
orient='v', ax=axes[0, 1])
```

```
axes[1,0].set_title("Distribution of Petal Length")

sns.boxplot(y="petal_length", x="species", data=dataset,
orient='v', ax=axes[1, 0])
```

```
axes[1,1].set_title("Distribution of Petal Width")

sns.boxplot(y="petal_width", x="species", data=dataset,
orient='v', ax=axes[1, 1])
```

```
plt.show()
```

- **sns.boxplot()**: This function is used to create a boxplot for each feature in the dataset, grouped by species (**x="species"**).
- **orient='v'**: The **orient** parameter specifies the orientation of the box plot. 'v' means vertical, and 'h' would mean horizontal.
- **ax=axes[x, y]**: This specifies the subplot in which the boxplot is drawn.
- These boxplots show the distribution of each feature for each Iris species. They help us identify the median, quartiles, and outliers in the data.

## 6. Inferences

The dataset consists of four features (numeric) and one target variable **species** (nominal). Here are the results based on the analysis:

### 1. List of Features and Their Types:

- **sepal\_length**: Numeric
- **sepal\_width**: Numeric
- **petal\_length**: Numeric
- **petal\_width**: Numeric
- **species**: Nominal (Categorical)

### 2. Histograms:

- **Sepal Length**: The distribution is relatively symmetric, with a higher frequency around 5.0 to 6.0.
- **Sepal Width**: The distribution is slightly skewed with a peak around 3.0.
- **Petal Length**: There is a peak around 1.5 to 5.0, with some longer petals observed, especially for the "virginica" species.
- **Petal Width**: Similar to petal length, petal width has a peak around 0.2-1.0, with wider petals in "virginica."

### 3. Boxplots:

- The boxplots show that the "versicolor" species has a similar sepal length to "virginica," but "setosa" has a distinctively smaller range.
- "Virginica" has the largest petal length and width, with a few outliers in both the petal and sepal features.
- "Setosa" tends to have the smallest values for petal length and width compared to the other species.

### 4. Outliers:

- **The boxplots reveal a few outliers, especially for "virginica," where petal length and width have some extreme values compared to the rest of the data.**