

```
In [3]: import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import warnings
warnings.simplefilter('ignore')
df=pd.read_csv(r"C:\Users\HP\Downloads\New folder\1) iris (1).csv")
df
```

```
Out[3]:
```

|     | 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    |
| ... | ...          | ...         | ...          | ...         | ...       |
| 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 |

150 rows × 5 columns

```
In [4]: df.isnull()
```

Out[4]:

|     | sepal_length | sepal_width | petal_length | petal_width | species |
|-----|--------------|-------------|--------------|-------------|---------|
| 0   | False        | False       | False        | False       | False   |
| 1   | False        | False       | False        | False       | False   |
| 2   | False        | False       | False        | False       | False   |
| 3   | False        | False       | False        | False       | False   |
| 4   | False        | False       | False        | False       | False   |
| ... | ...          | ...         | ...          | ...         | ...     |
| 145 | False        | False       | False        | False       | False   |
| 146 | False        | False       | False        | False       | False   |
| 147 | False        | False       | False        | False       | False   |
| 148 | False        | False       | False        | False       | False   |
| 149 | False        | False       | False        | False       | False   |

150 rows × 5 columns

In [5]: `df.describe()`

Out[5]:

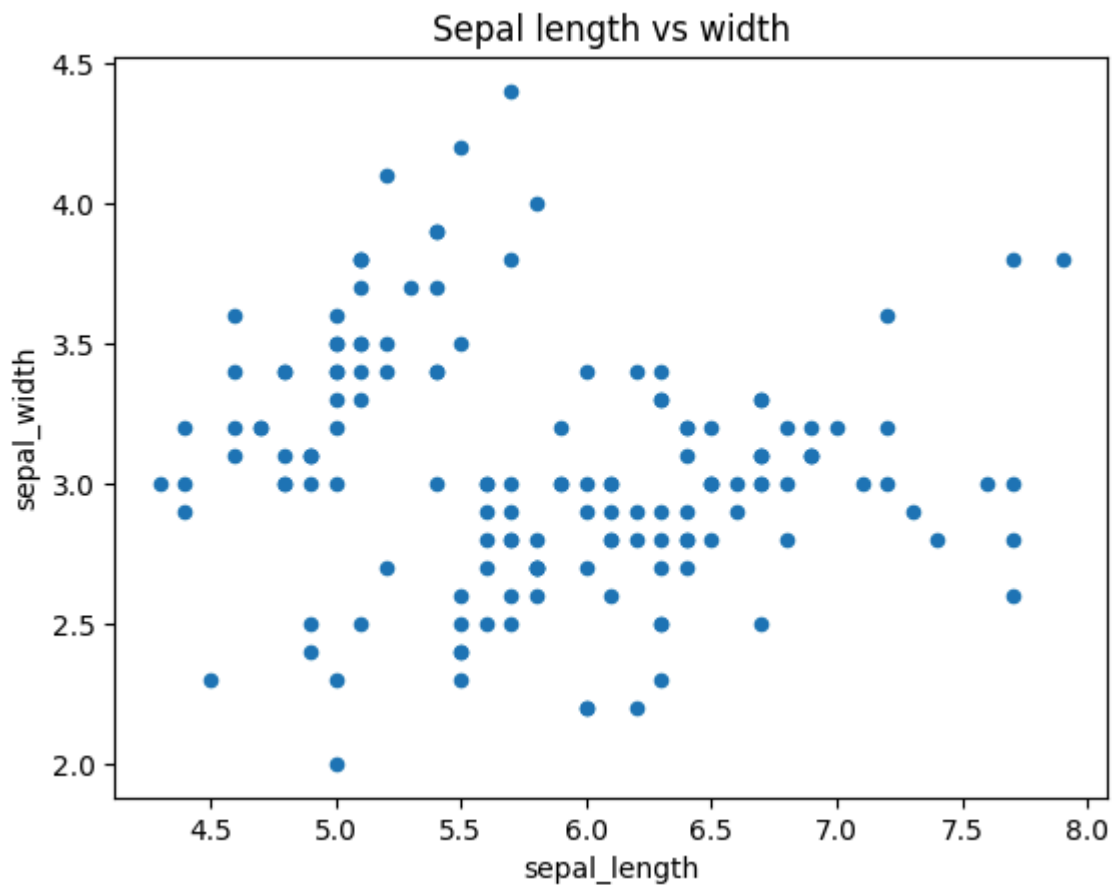
|       | sepal_length | sepal_width | petal_length | petal_width |
|-------|--------------|-------------|--------------|-------------|
| count | 150.000000   | 150.000000  | 150.000000   | 150.000000  |
| mean  | 5.843333     | 3.054000    | 3.758667     | 1.198667    |
| std   | 0.828066     | 0.433594    | 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    |

In [6]: `df.isnull().sum()`

Out[6]:

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

In [7]: `df.plot(kind="scatter", x="sepal_length", y="sepal_width", title="Sepal length v  
plt.show()`



```
In [8]: #Load the dataset:
```

```
In [9]: import seaborn as sns
import matplotlib.pyplot as plt

# Load iris dataset (seaborn has it built-in)
df = sns.load_dataset("iris")

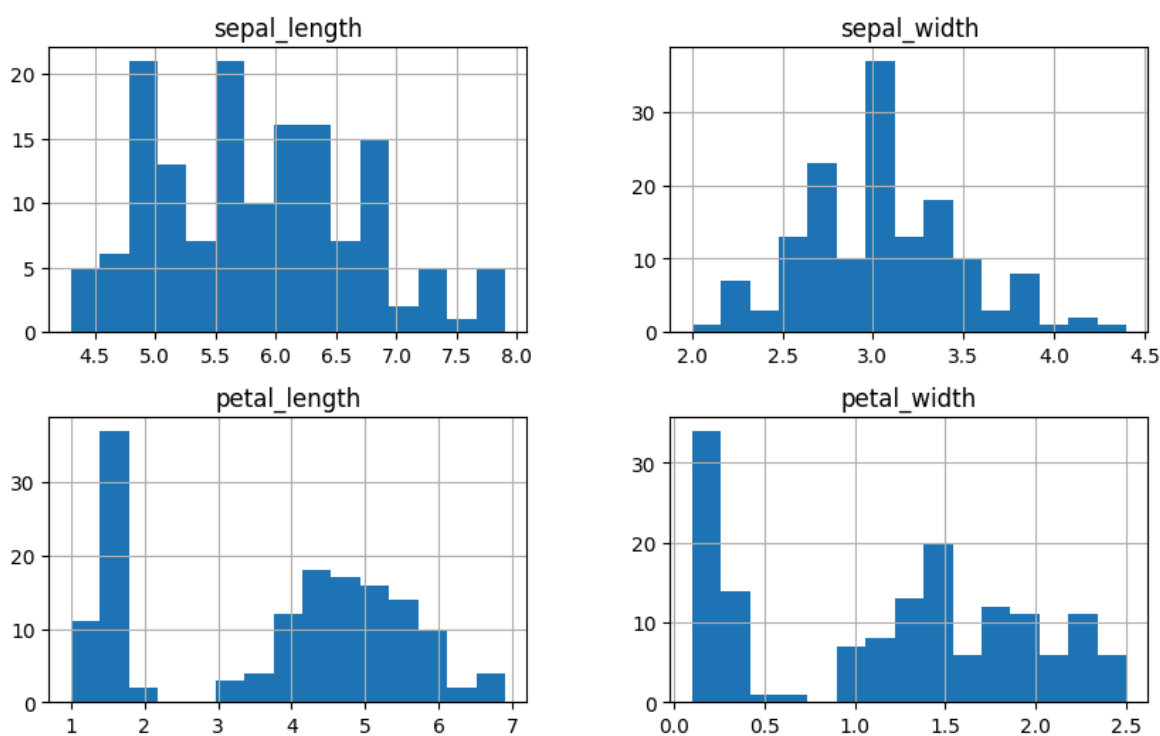
print(df.head())
```

|   | 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  |

```
In [10]: #plot using histogram:
```

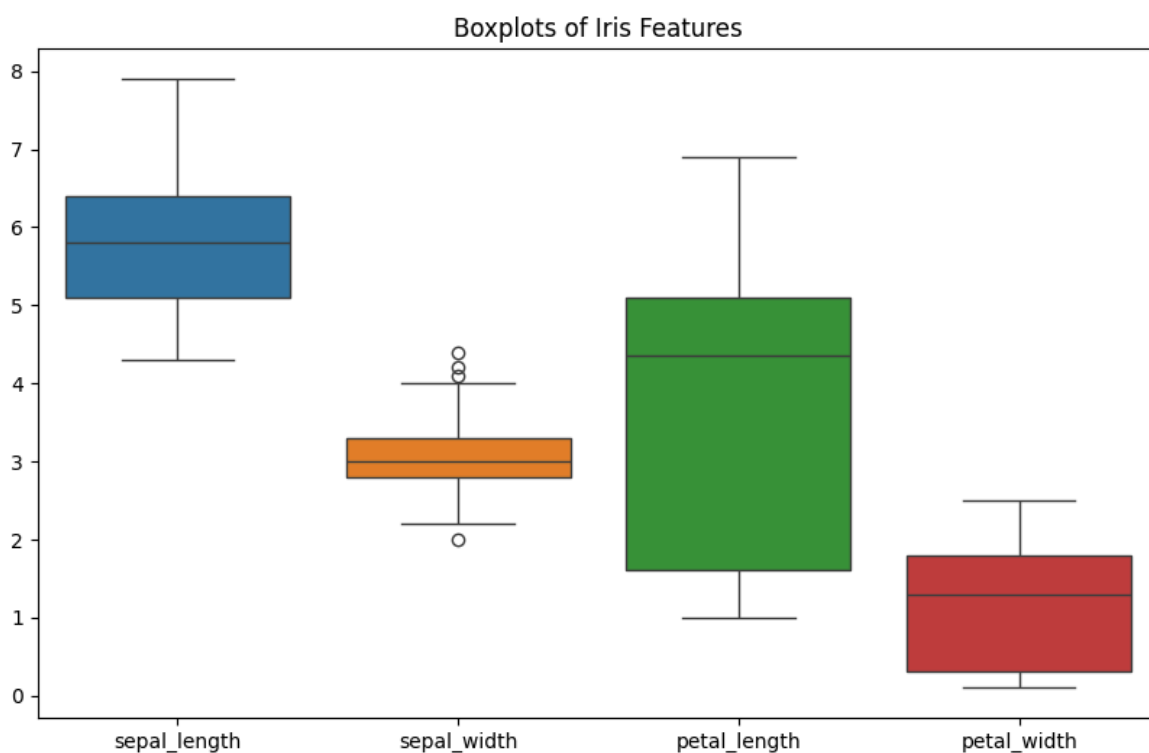
```
In [11]: df.hist(figsize=(10, 6), bins=15)
plt.suptitle("Feature Distributions", fontsize=16)
plt.show()
```

## Feature Distributions



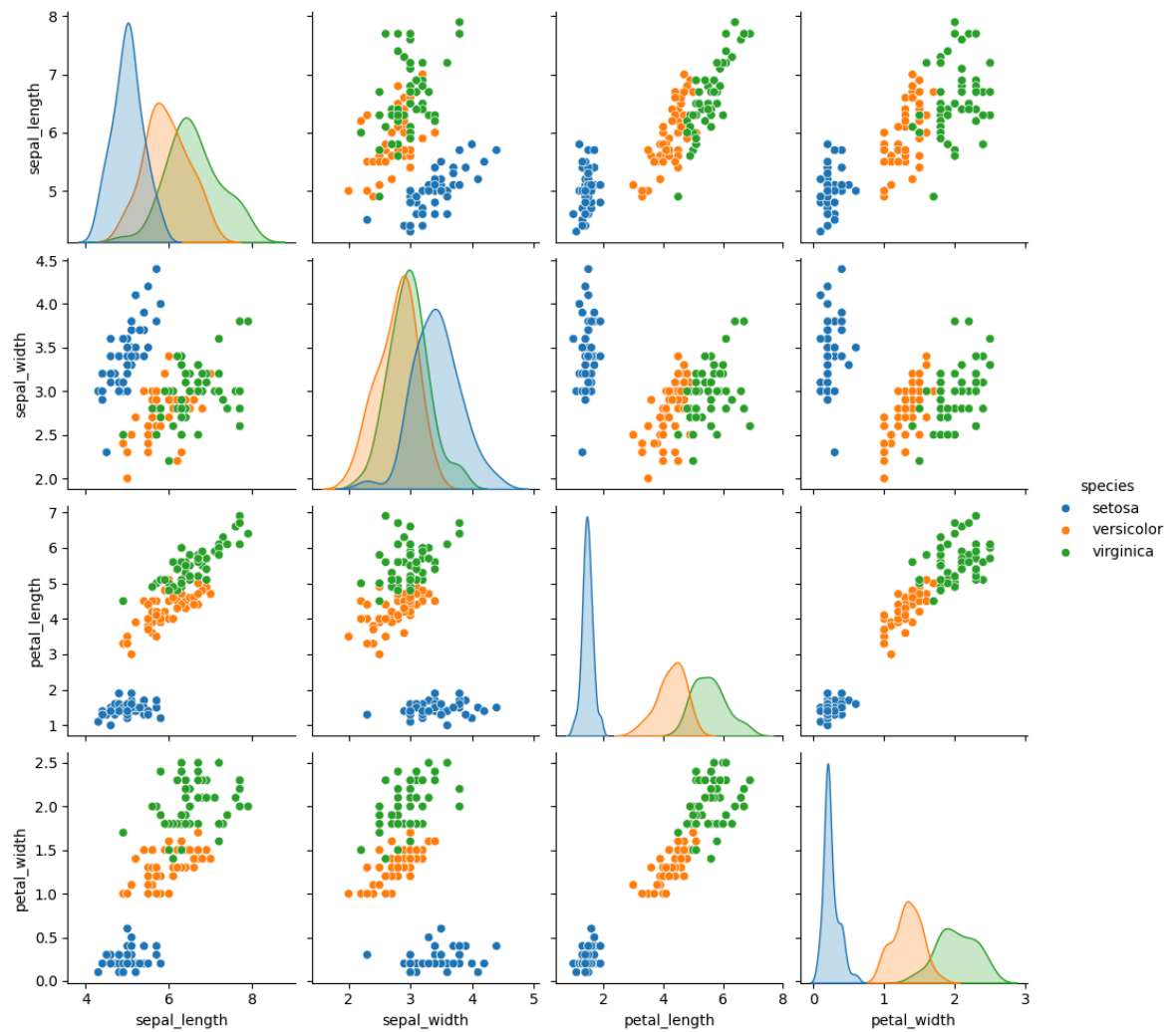
In [12]: *#Boxplots - Detect spread & outliers:*

```
In [13]: plt.figure(figsize=(10, 6))
sns.boxplot(data=df)
plt.title("Boxplots of Iris Features")
plt.show()
```



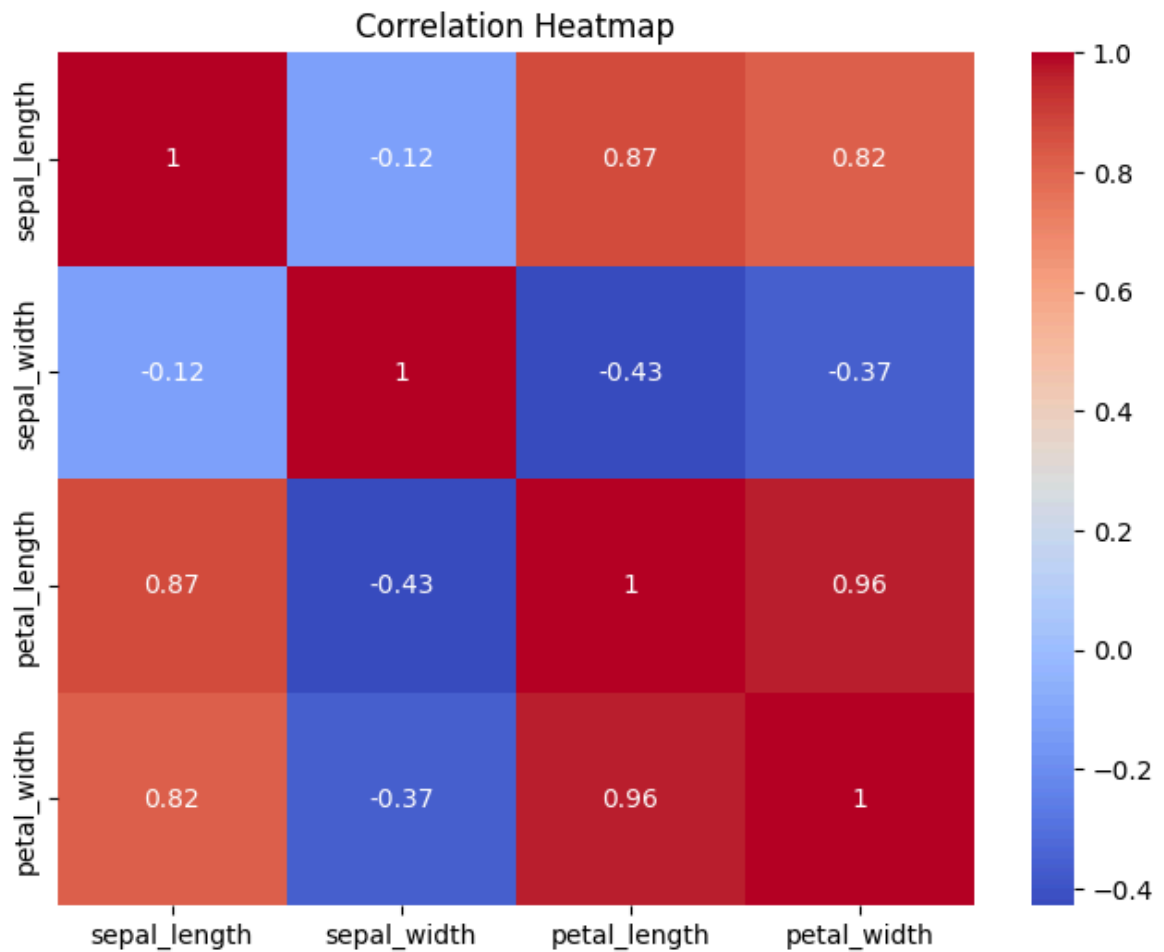
In [14]: *#Pairplot - Relationships between all features:*

```
In [15]: sns.pairplot(df, hue="species", diag_kind="kde")
plt.show()
```



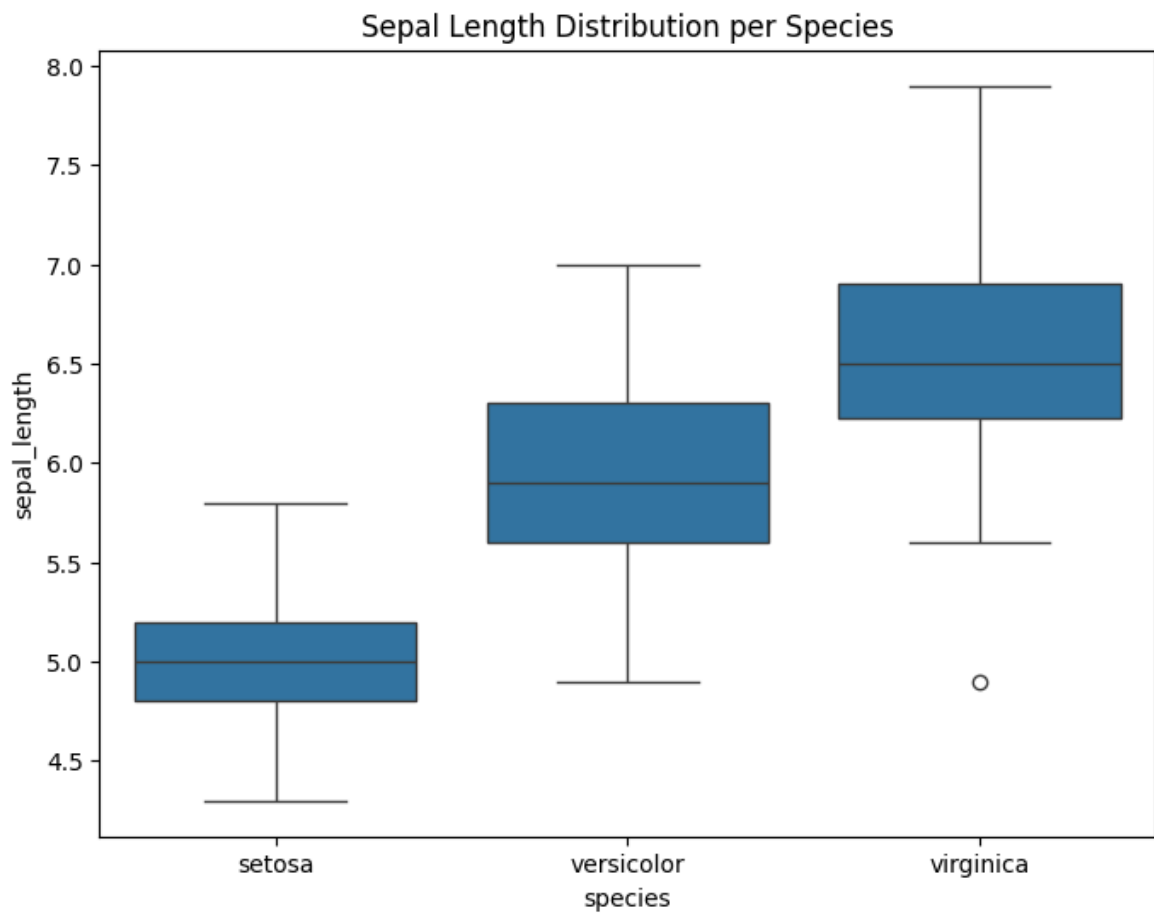
In [16]: *#Correlation Heatmap:*

```
In [17]: plt.figure(figsize=(8, 6))
sns.heatmap(df.drop("species", axis=1).corr(), annot=True, cmap="coolwarm")
plt.title("Correlation Heatmap")
plt.show()
```



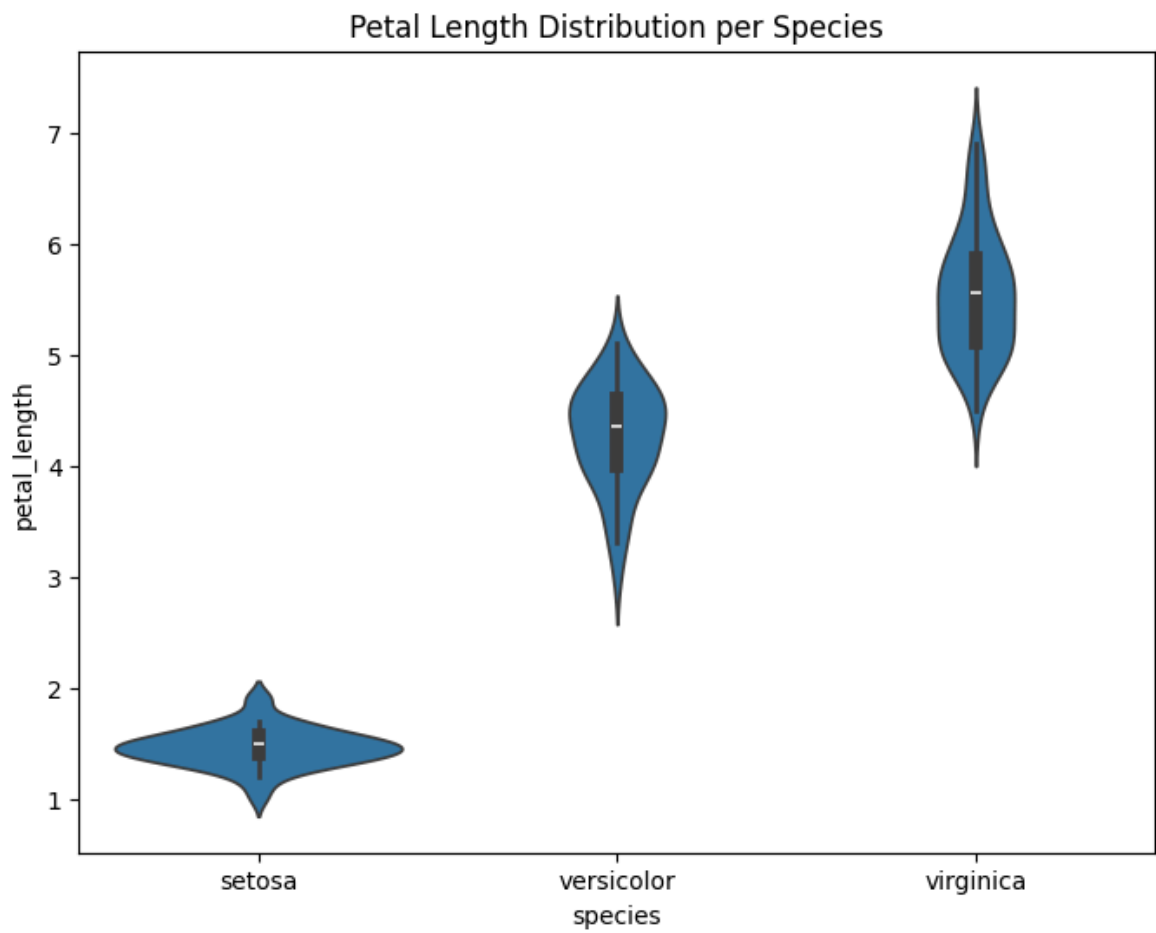
```
In [18]: #Species-wise Boxplot (Feature Comparison):
```

```
In [19]: plt.figure(figsize=(8, 6))  
sns.boxplot(x="species", y="sepal_length", data=df)  
plt.title("Sepal Length Distribution per Species")  
plt.show()
```



In [20]: *#Violin Plots - Feature Distribution by Species:*

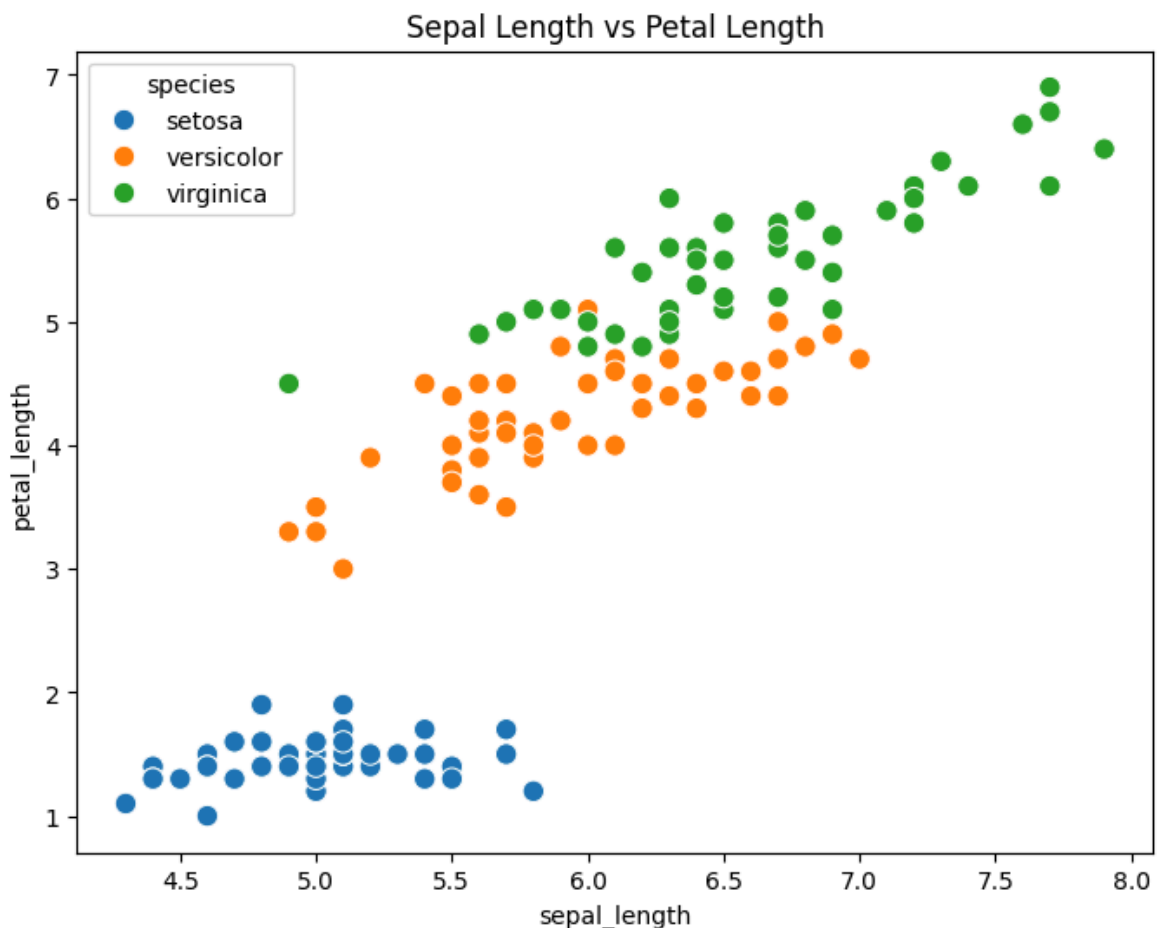
```
In [21]: plt.figure(figsize=(8, 6))
sns.violinplot(x="species", y="petal_length", data=df)
plt.title("Petal Length Distribution per Species")
plt.show()
```



In [22]: *#Scatter Plot (2D Feature Space):*

```
In [23]: plt.figure(figsize=(8, 6))
sns.scatterplot(x="sepal_length", y="petal_length", hue="species", data=df, s=80)
plt.title("Sepal Length vs Petal Length")
plt.show()
```





```
In [24]: # Dashboard-style visualization of Iris dataset
import seaborn as sns
import matplotlib.pyplot as plt
import pandas as pd

# Load dataset
df = sns.load_dataset("iris")

# Set style
sns.set(style="whitegrid")

# Create figure and subplots
fig, axes = plt.subplots(2, 3, figsize=(18, 10))

# 1. Histogram
df.drop("species", axis=1).hist(ax=axes[0,0], bins=10, color="skyblue")
axes[0,0].set_title("Feature Distributions")

# 2. Boxplot by feature
sns.boxplot(data=df.drop("species", axis=1), ax=axes[0,1])
axes[0,1].set_title("Boxplots of Features")

# 3. Heatmap
sns.heatmap(df.drop("species", axis=1).corr(), annot=True, cmap="coolwarm", ax=axes[0,2])
axes[0,2].set_title("Correlation Heatmap")

# 4. Scatterplot
sns.scatterplot(x="sepal_length", y="petal_length", hue="species", data=df, ax=axes[1,0])
axes[1,0].set_title("Sepal Length vs Petal Length")

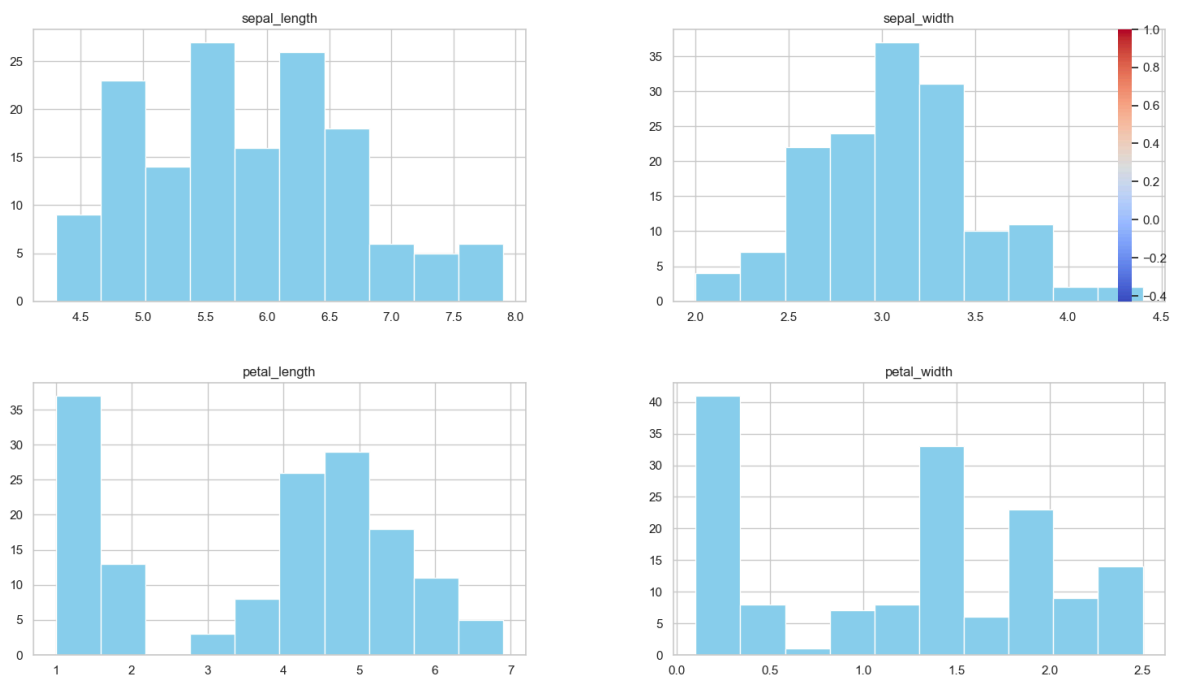
# 5. Boxplot by species
```

```
sns.boxplot(x="species", y="sepal_length", data=df, ax=axes[1,1])
axes[1,1].set_title("Sepal Length by Species")

# 6. Violin plot
sns.violinplot(x="species", y="petal_length", data=df, ax=axes[1,2])
axes[1,2].set_title("Petal Length by Species")

plt.suptitle("Iris Dataset Dashboard", fontsize=18, fontweight="bold")
plt.tight_layout(rect=[0, 0, 1, 0.96]) # Adjust Layout
plt.show()
```

Iris Dataset Dashboard



```
In [25]: # Dashboard + Pairplot visualization of Iris dataset
import seaborn as sns
import matplotlib.pyplot as plt

# Load dataset
df = sns.load_dataset("iris")

# Set style
sns.set(style="whitegrid")

# ----- Dashboard -----
fig, axes = plt.subplots(2, 3, figsize=(18, 10))

# 1. Histogram
df.drop("species", axis=1).hist(ax=axes[0,0], bins=10, color="skyblue")
axes[0,0].set_title("Feature Distributions")

# 2. Boxplot by feature
sns.boxplot(data=df.drop("species", axis=1), ax=axes[0,1])
axes[0,1].set_title("Boxplots of Features")

# 3. Heatmap
sns.heatmap(df.drop("species", axis=1).corr(), annot=True, cmap="coolwarm", ax=axes[0,2])
axes[0,2].set_title("Correlation Heatmap")

# 4. Scatterplot
```

```

sns.scatterplot(x="sepal_length", y="petal_length", hue="species", data=df, ax=axes[1,0])
axes[1,0].set_title("Sepal Length vs Petal Length")

# 5. Boxplot by species
sns.boxplot(x="species", y="sepal_length", data=df, ax=axes[1,1])
axes[1,1].set_title("Sepal Length by Species")

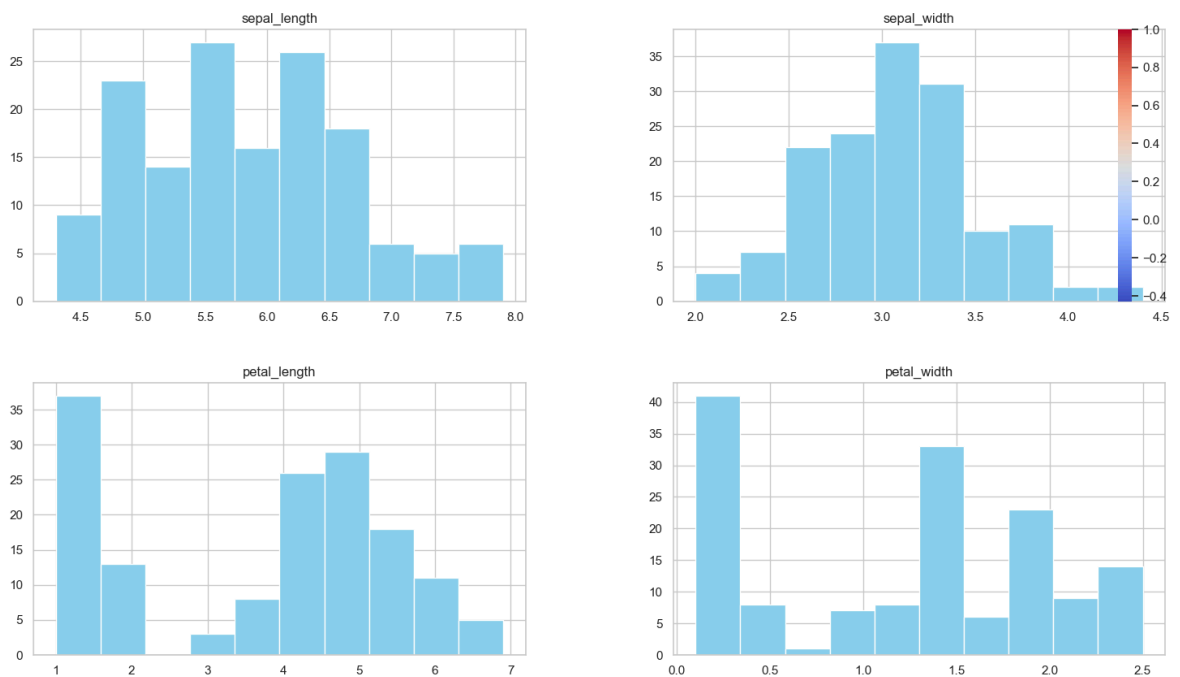
# 6. Violin plot
sns.violinplot(x="species", y="petal_length", data=df, ax=axes[1,2])
axes[1,2].set_title("Petal Length by Species")

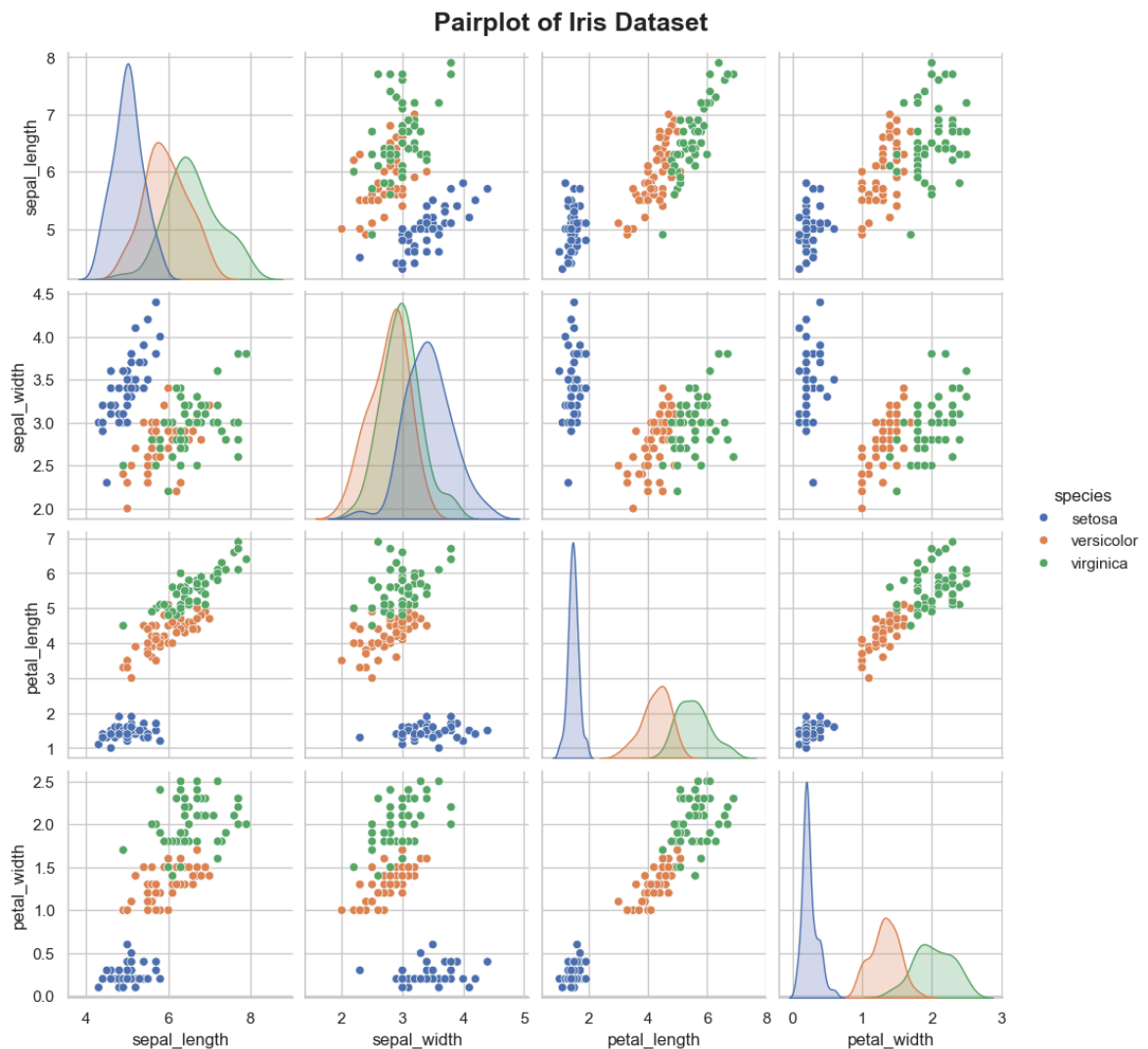
plt.suptitle("Iris Dataset Dashboard", fontsize=18, fontweight="bold")
plt.tight_layout(rect=[0, 0, 1, 0.96])
plt.show()

# ----- Pairplot -----
sns.pairplot(df, hue="species", diag_kind="kde")
plt.suptitle("Pairplot of Iris Dataset", fontsize=18, fontweight="bold", y=1.02)
plt.show()

```

Iris Dataset Dashboard





```
In [26]: # Iris Dataset: Dashboard + Pairplot + 3D Scatter
import seaborn as sns
import matplotlib.pyplot as plt
from mpl_toolkits.mplot3d import Axes3D # for 3D plots

# Load dataset
df = sns.load_dataset("iris")
sns.set(style="whitegrid")

# ----- Dashboard -----
fig, axes = plt.subplots(2, 3, figsize=(18, 10))

# 1. Histogram
df.drop("species", axis=1).hist(ax=axes[0,0], bins=10, color="skyblue")
axes[0,0].set_title("Feature Distributions")

# 2. Boxplot by feature
sns.boxplot(data=df.drop("species", axis=1), ax=axes[0,1])
axes[0,1].set_title("Boxplots of Features")

# 3. Heatmap
sns.heatmap(df.drop("species", axis=1).corr(), annot=True, cmap="coolwarm", ax=axes[0,2])
axes[0,2].set_title("Correlation Heatmap")

# 4. Scatterplot
sns.scatterplot(x="sepal_length", y="petal_length", hue="species", data=df, ax=axes[1,0])
axes[1,0].set_title("Sepal Length vs Petal Length")
```

```
# 5. Boxplot by species
sns.boxplot(x="species", y="sepal_length", data=df, ax=axes[1,1])
axes[1,1].set_title("Sepal Length by Species")

# 6. Violin plot
sns.violinplot(x="species", y="petal_length", data=df, ax=axes[1,2])
axes[1,2].set_title("Petal Length by Species")

plt.suptitle("Iris Dataset Dashboard", fontsize=18, fontweight="bold")
plt.tight_layout(rect=[0, 0, 1, 0.96])
plt.show()

# ----- Pairplot -----
sns.pairplot(df, hue="species", diag_kind="kde")
plt.suptitle("Pairplot of Iris Dataset", fontsize=18, fontweight="bold", y=1.02)
plt.show()

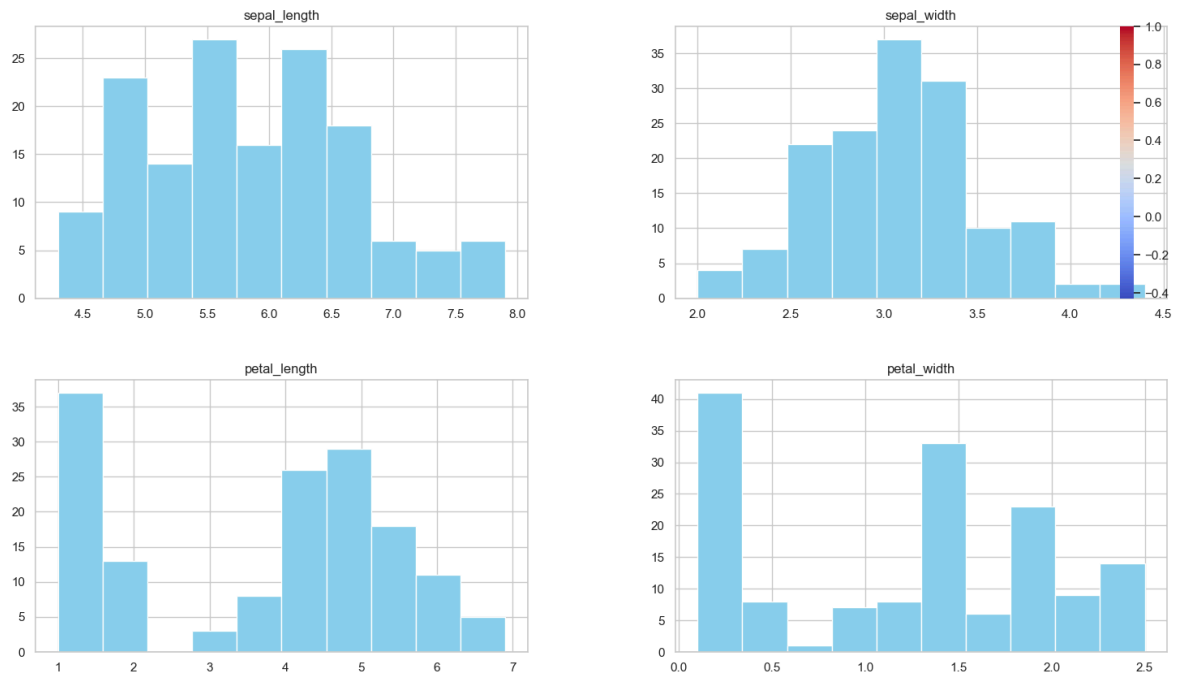
# ----- 3D Scatter Plot -----
fig = plt.figure(figsize=(10, 7))
ax = fig.add_subplot(111, projection="3d")

species_list = df["species"].unique()
colors = ["red", "green", "blue"]

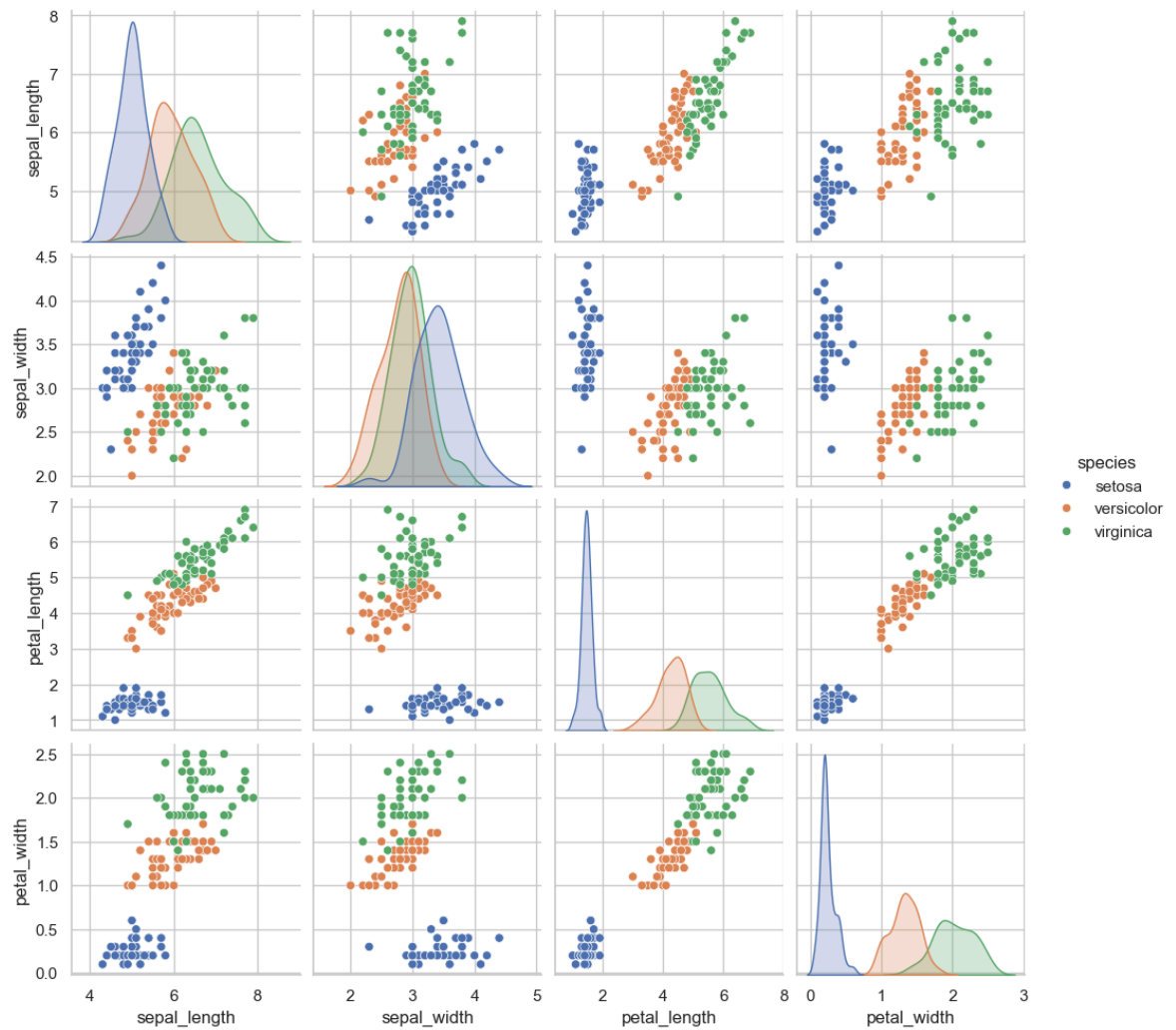
for species, color in zip(species_list, colors):
    subset = df[df["species"] == species]
    ax.scatter(
        subset["sepal_length"],
        subset["sepal_width"],
        subset["petal_length"],
        c=color, label=species, s=60
    )

ax.set_xlabel("Sepal Length")
ax.set_ylabel("Sepal Width")
ax.set_zlabel("Petal Length")
ax.set_title("3D Scatter Plot of Iris Dataset")
ax.legend()
plt.show()
```

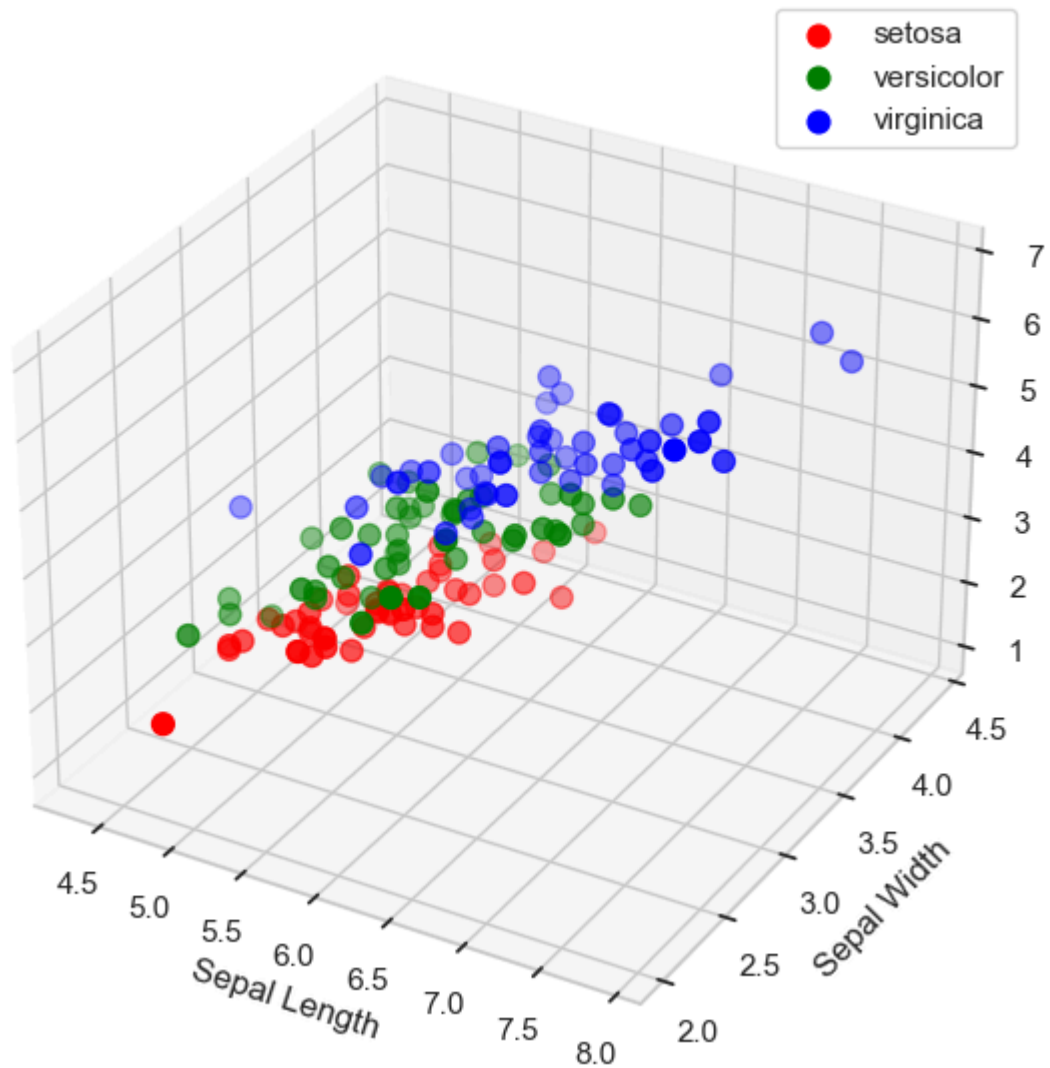
## Iris Dataset Dashboard



## Pairplot of Iris Dataset



3D Scatter Plot of Iris Dataset



```
In [27]: # Iris Dataset: Dashboard + Pairplot + 3D + 4D Scatter
import seaborn as sns
import matplotlib.pyplot as plt
from mpl_toolkits.mplot3d import Axes3D

# Load dataset
df = sns.load_dataset("iris")
sns.set(style="whitegrid")

# ----- Dashboard -----
fig, axes = plt.subplots(2, 3, figsize=(18, 10))

# 1. Histogram
df.drop("species", axis=1).hist(ax=axes[0,0], bins=10, color="skyblue")
axes[0,0].set_title("Feature Distributions")

# 2. Boxplot by feature
sns.boxplot(data=df.drop("species", axis=1), ax=axes[0,1])
axes[0,1].set_title("Boxplots of Features")

# 3. Heatmap
sns.heatmap(df.drop("species", axis=1).corr(), annot=True, cmap="coolwarm", ax=axes[0,2])
axes[0,2].set_title("Correlation Heatmap")

# 4. Scatterplot
```

```

sns.scatterplot(x="sepal_length", y="petal_length", hue="species", data=df, ax=axes[1,0]).set_title("Sepal Length vs Petal Length")

# 5. Boxplot by species
sns.boxplot(x="species", y="sepal_length", data=df, ax=axes[1,1])
axes[1,1].set_title("Sepal Length by Species")

# 6. Violin plot
sns.violinplot(x="species", y="petal_length", data=df, ax=axes[1,2])
axes[1,2].set_title("Petal Length by Species")

plt.suptitle("Iris Dataset Dashboard", fontsize=18, fontweight="bold")
plt.tight_layout(rect=[0, 0, 1, 0.96])
plt.show()

# ----- Pairplot -----
sns.pairplot(df, hue="species", diag_kind="kde")
plt.suptitle("Pairplot of Iris Dataset", fontsize=18, fontweight="bold", y=1.02)
plt.show()

# ----- 3D Scatter Plot -----
fig = plt.figure(figsize=(10, 7))
ax = fig.add_subplot(111, projection="3d")

species_list = df["species"].unique()
colors = ["red", "green", "blue"]

for species, color in zip(species_list, colors):
    subset = df[df["species"] == species]
    ax.scatter(
        subset["sepal_length"],
        subset["sepal_width"],
        subset["petal_length"],
        c=color, label=species, s=60
    )

ax.set_xlabel("Sepal Length")
ax.set_ylabel("Sepal Width")
ax.set_zlabel("Petal Length")
ax.set_title("3D Scatter Plot of Iris Dataset")
ax.legend()
plt.show()

# ----- 4D Scatter Plot -----
fig = plt.figure(figsize=(10, 7))
ax = fig.add_subplot(111, projection="3d")

for species, color in zip(species_list, colors):
    subset = df[df["species"] == species]
    ax.scatter(
        subset["sepal_length"],
        subset["sepal_width"],
        subset["petal_length"],
        c=color, s=subset["petal_width"] * 40, # size = petal_width
        alpha=0.7, label=species
    )

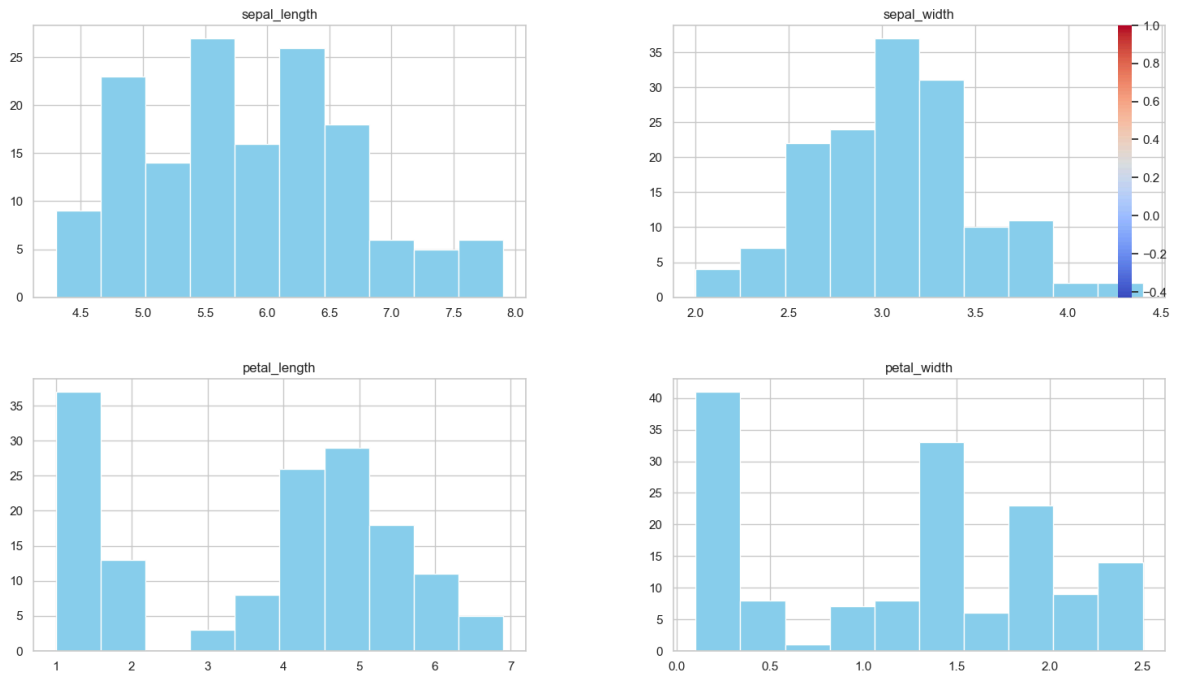
ax.set_xlabel("Sepal Length")
ax.set_ylabel("Sepal Width")
ax.set_zlabel("Petal Length")

```

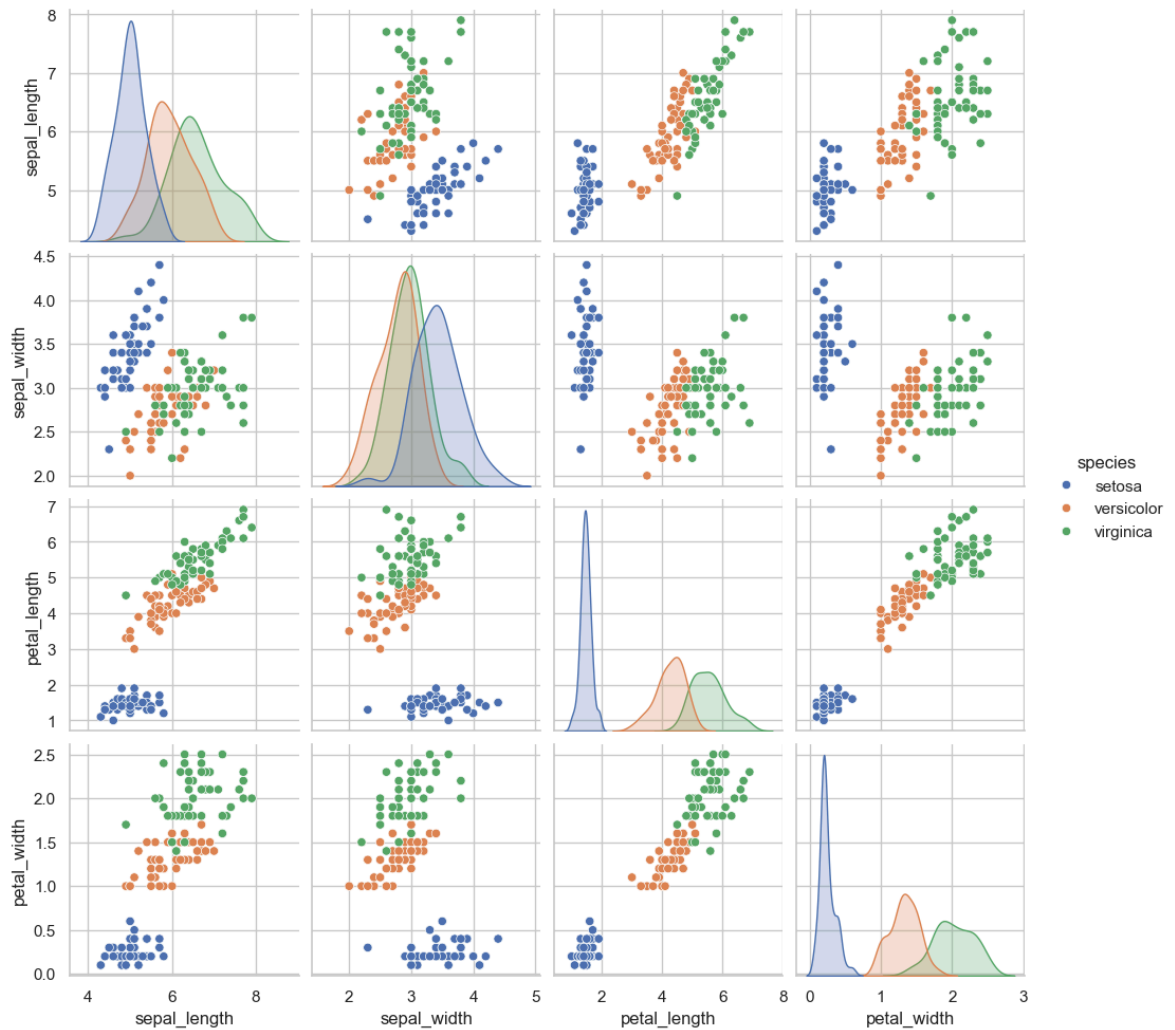


```
ax.set_title("4D Scatter Plot (Petal Width as Size)")
ax.legend()
plt.show()
```

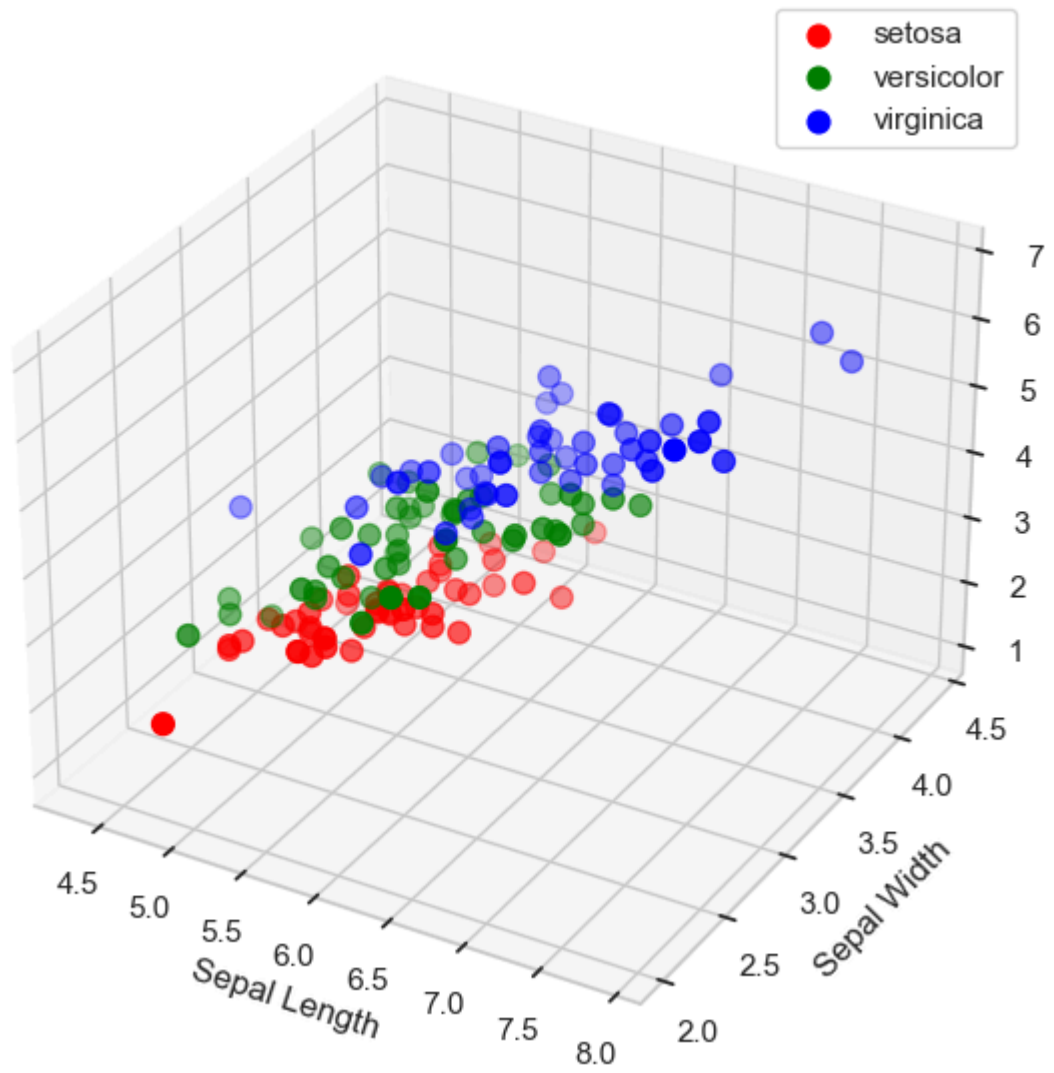
Iris Dataset Dashboard



Pairplot of Iris Dataset



3D Scatter Plot of Iris Dataset



4D Scatter Plot (Petal Width as Size)

