

```

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

iris = sns.load_dataset('iris')
iris.to_csv('IRIS(1).csv', index=False)

iris.info()

<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

iris.head()

{"summary": "{\n  \"name\": \"iris\", \n  \"rows\": 150, \n  \"fields\": [\n    {\n      \"column\": \"sepal_length\", \n      \"properties\": {\n        \"dtype\": \"number\", \n        \"std\": 0.8280661279778629, \n        \"min\": 4.3, \n        \"max\": 7.9, \n        \"num_unique_values\": 35, \n        \"samples\": [\n          6.2, \n          4.5, \n          5.6\n        ], \n        \"semantic_type\": \"\", \n        \"description\": \"\", \n        \"column\": \"sepal_width\", \n        \"properties\": {\n          \"dtype\": \"number\", \n          \"std\": 0.435866284936698, \n          \"min\": 2.0, \n          \"max\": 4.4, \n          \"num_unique_values\": 23, \n          \"samples\": [\n            2.3, \n            4.0, \n            3.5\n          ], \n          \"semantic_type\": \"\", \n          \"description\": \"\", \n          \"column\": \"petal_length\", \n          \"properties\": {\n            \"dtype\": \"number\", \n            \"std\": 1.7652982332594667, \n            \"min\": 1.0, \n            \"max\": 6.9, \n            \"num_unique_values\": 43, \n            \"samples\": [\n              6.7, \n              3.8, \n              3.7\n            ], \n            \"semantic_type\": \"\", \n            \"description\": \"\", \n            \"column\": \"petal_width\", \n            \"properties\": {\n              \"dtype\": \"number\", \n              \"std\": 0.7622376689603465, \n              \"min\": 0.1, \n              \"max\": 2.5, \n              \"num_unique_values\": 22, \n              \"samples\": [\n                0.2, \n                1.2, \n                1.3\n              ], \n              \"semantic_type\": \"\", \n              \"description\": \"\", \n              \"column\": \"species\", \n              \"properties\": {\n                \"dtype\": \"category\", \n                \"num_unique_values\": 3, \n                \"samples\": [\n                  \"Iris-setosa\", \n                  \"Iris-versicolor\", \n                  \"Iris-virginica\"\n                ]\n              }\n            }\n          }\n        }\n      }\n    }\n  ]\n}
```

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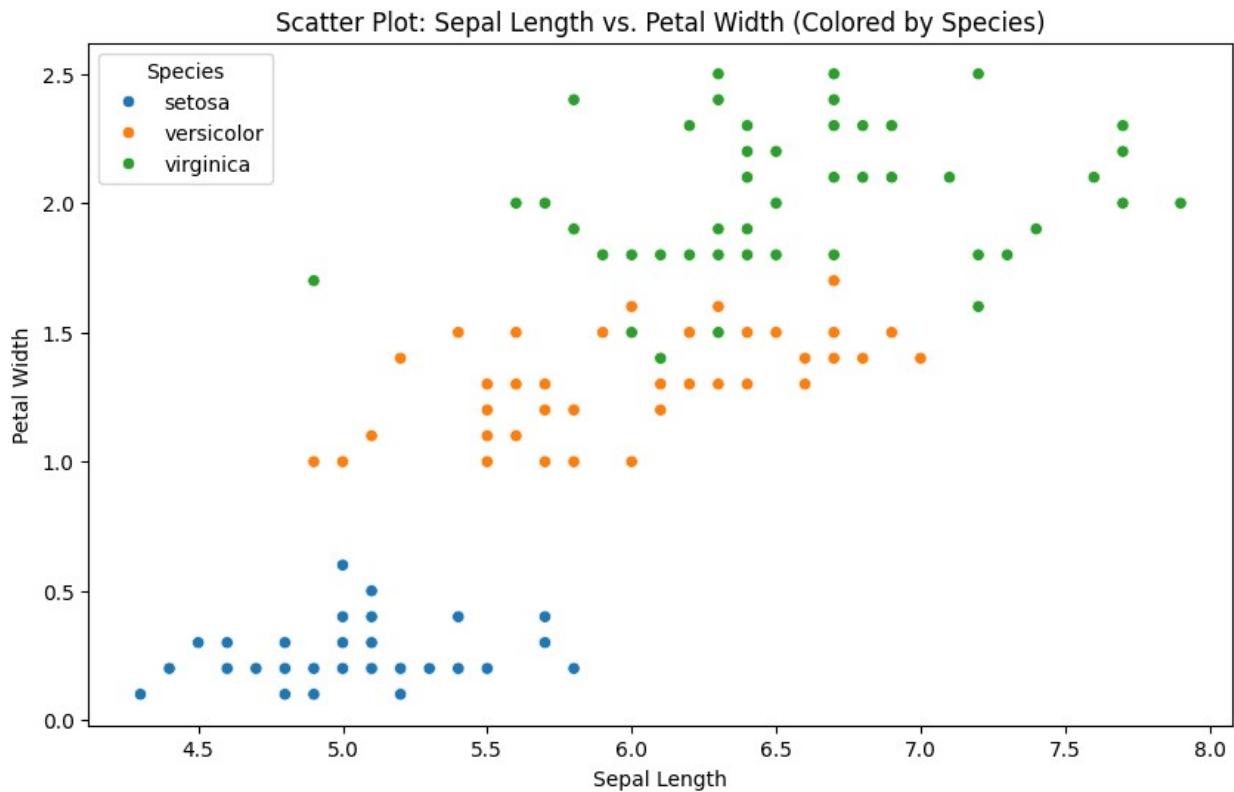
\"setosa\\"", \"versicolor\\"", \"virginica\"\n
],\n      \"semantic_type\": \"\", \n      \"description\": \"\"\n}\n  ]\n}, \"type\": \"dataframe\", \"variable_name\": \"iris\"}

iris.shape

(150, 5)

plt.figure(figsize=(10, 6))
sns.scatterplot(x='sepal_length', y='petal_width', hue='species',
data=iris)
plt.title('Scatter Plot: Sepal Length vs. Petal Width (Colored by Species)')
plt.xlabel('Sepal Length')
plt.ylabel('Petal Width')
plt.legend(title='Species')
plt.show()

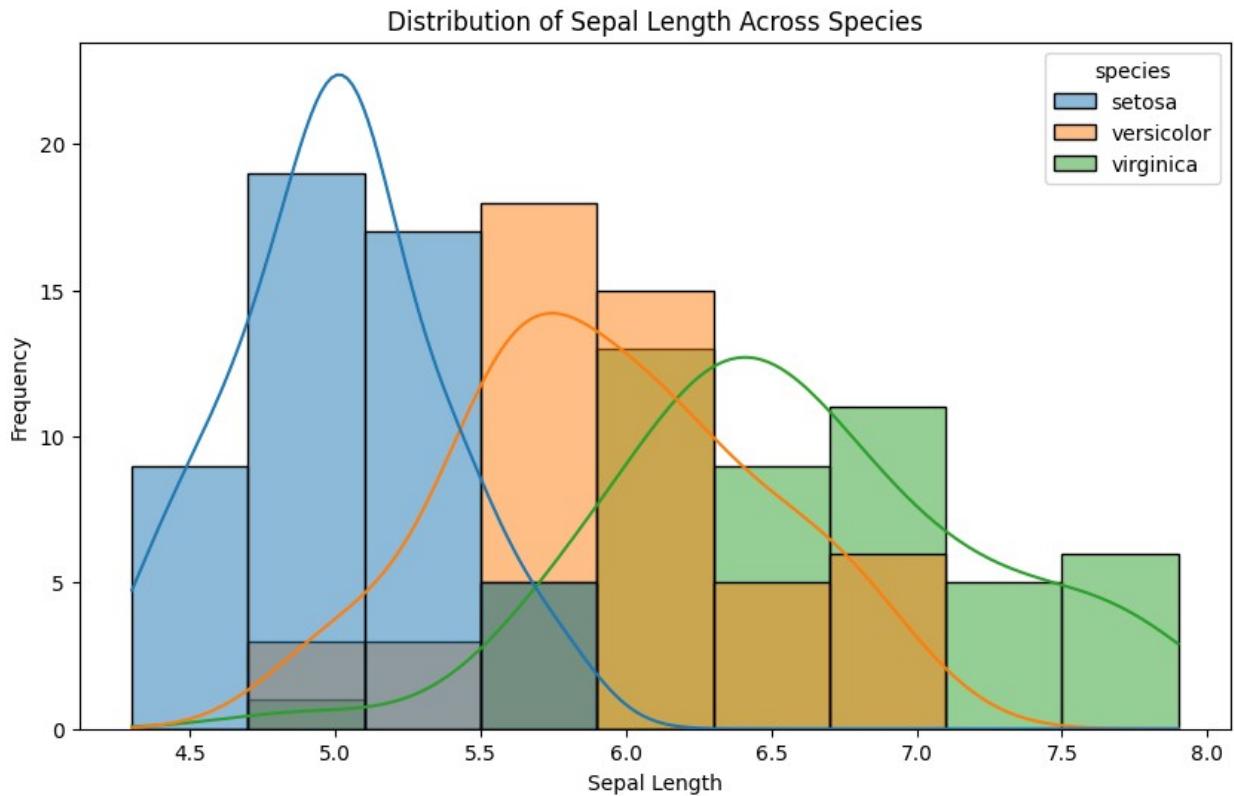
```



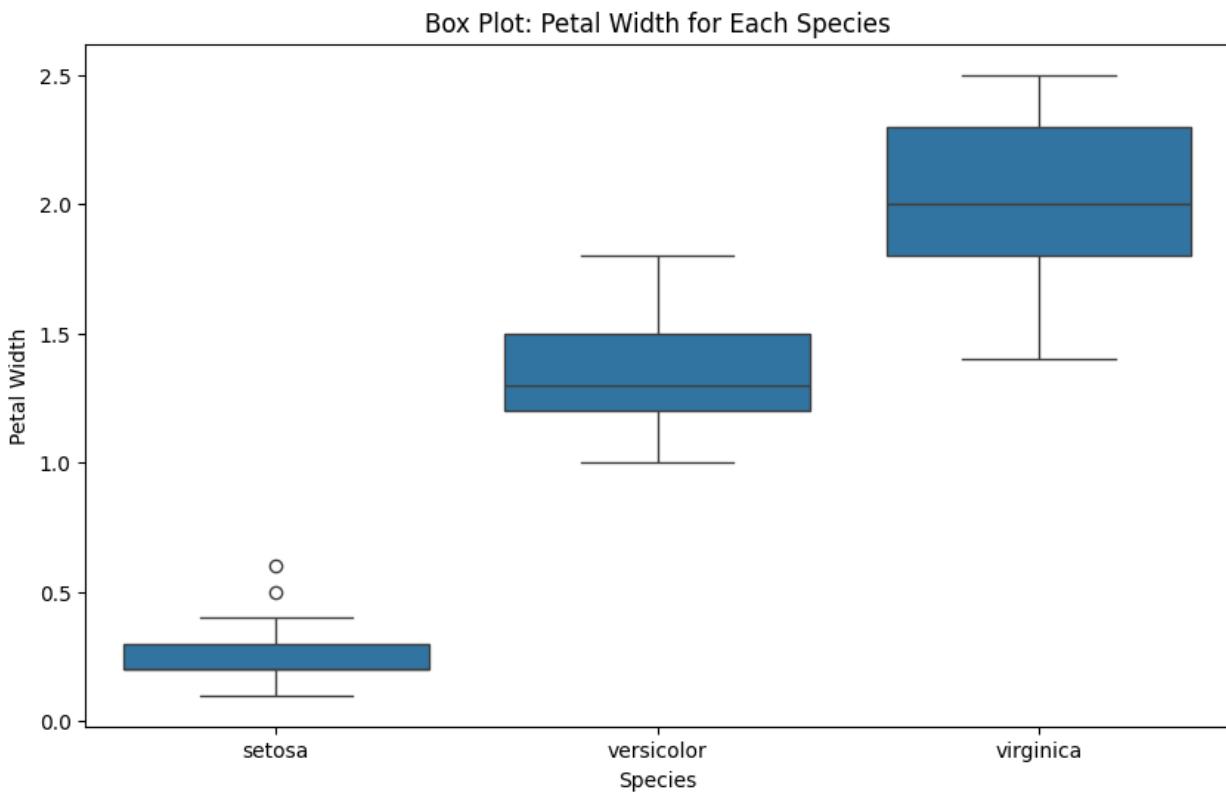
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plt.figure(figsize=(10, 6))
sns.histplot(data=iris, x='sepal_length', hue='species', kde=True)
plt.title('Distribution of Sepal Length Across Species')
plt.xlabel('Sepal Length')
plt.ylabel('Frequency')
plt.show()

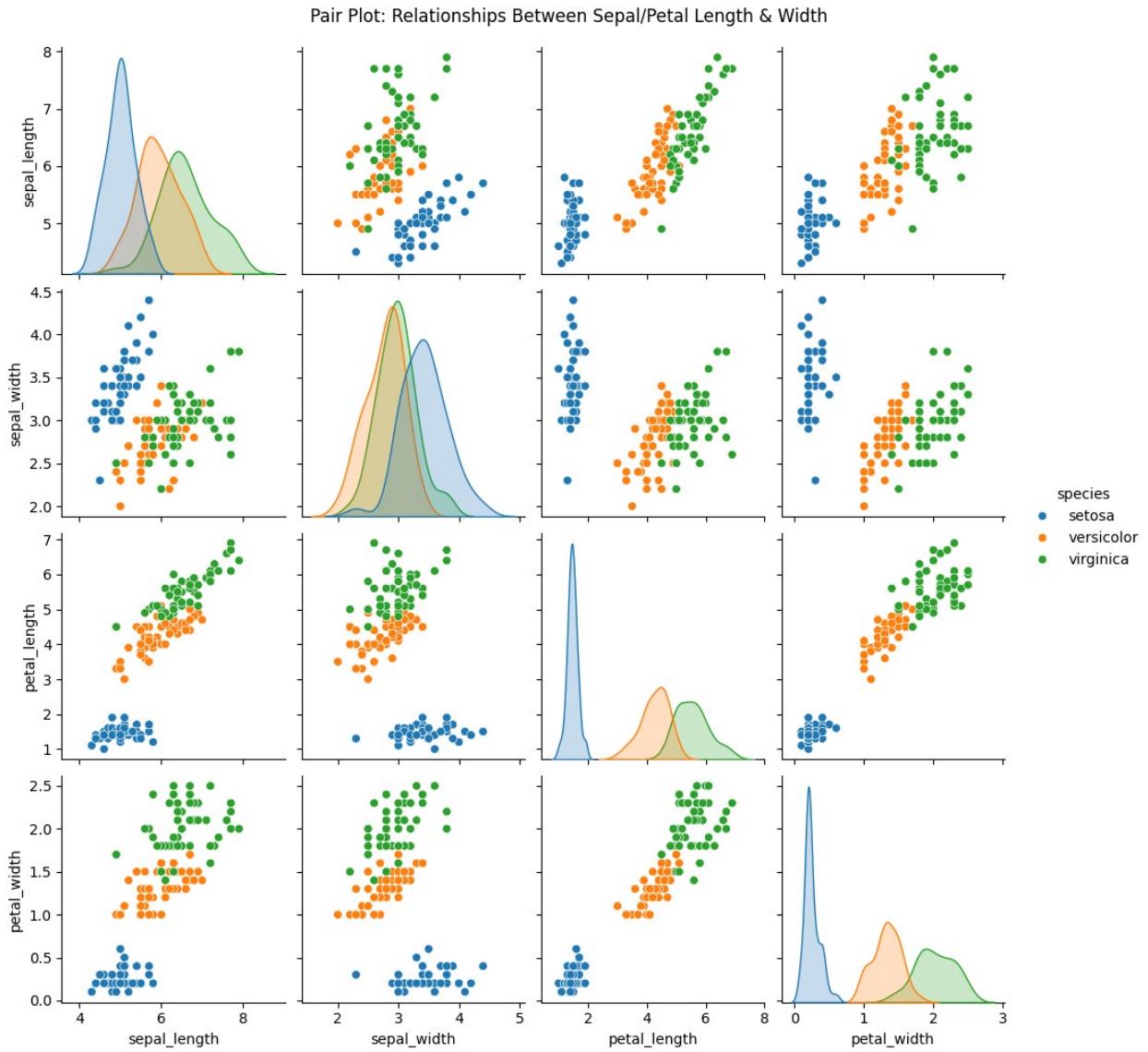
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```
plt.figure(figsize=(10, 6))
sns.boxplot(x='species', y='petal_width', data=iris)
plt.title('Box Plot: Petal Width for Each Species')
plt.xlabel('Species')
plt.ylabel('Petal Width')
plt.show()
```



```
sns.pairplot(iris, hue='species')
plt.suptitle('Pair Plot: Relationships Between Sepal/Petal Length &
Width', y=1.02)
plt.show()
```



```

plt.figure(figsize=(12, 8))
plt.subplot(2, 2, 1)
sns.violinplot(x='species', y='sepal_length', data=iris)
plt.title('Sepal Length by Species')

plt.subplot(2, 2, 2)
sns.violinplot(x='species', y='sepal_width', data=iris)
plt.title('Sepal Width by Species')

plt.subplot(2, 2, 3)
sns.violinplot(x='species', y='petal_length', data=iris)
plt.title('Petal Length by Species')

plt.subplot(2, 2, 4)

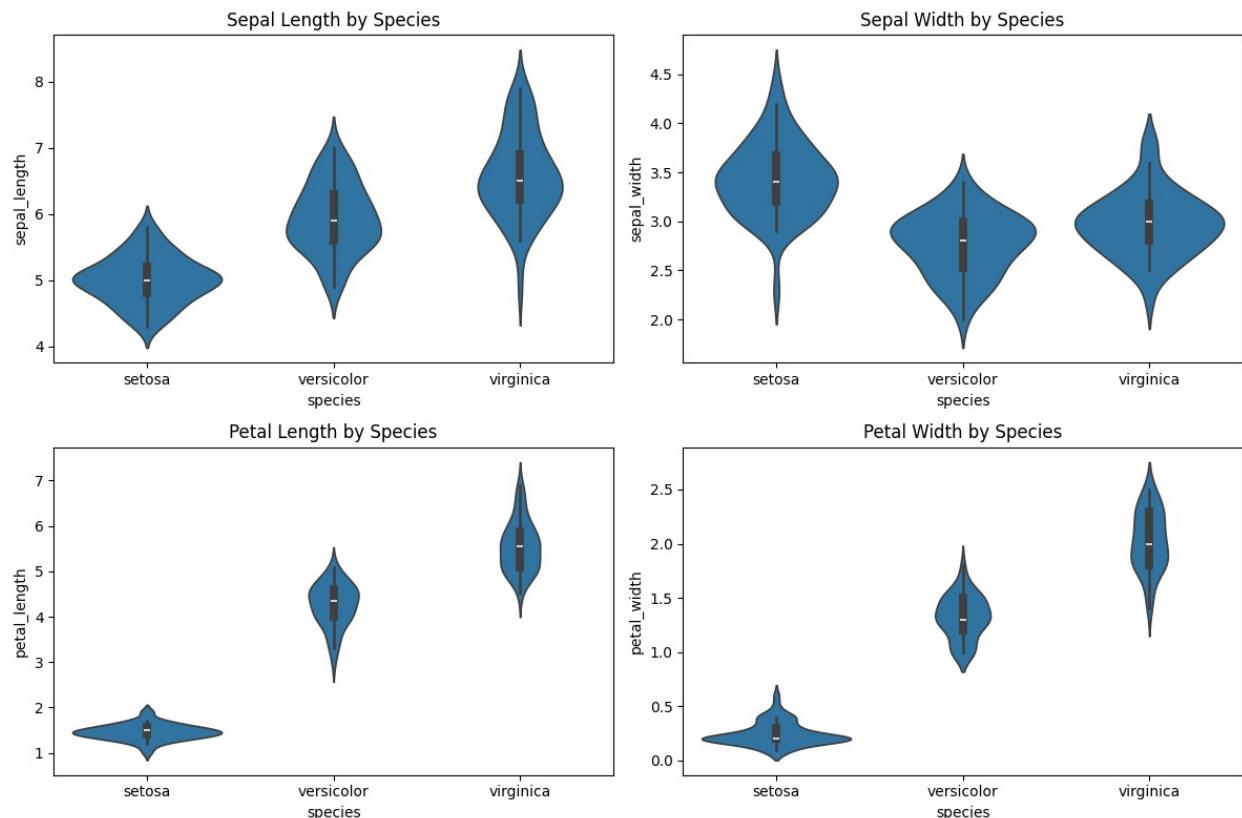
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sns.violinplot(x='species', y='petal_width', data=iris)
plt.title('Petal Width by Species')

plt.tight_layout()
plt.show()

```



```

plt.figure(figsize=(8, 6))
correlation_matrix = iris.corr(numeric_only=True)
sns.heatmap(correlation_matrix, annot=True, fmt='.2f')
plt.title('Correlation Matrix of Iris Dataset')
plt.show()

```

Correlation Matrix of Iris Dataset

