

```
In [12]: import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import LabelEncoder
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import accuracy_score, classification_report, confusion_matrix
```

```
In [4]: # loading the Dataset
IRIS_data = pd.read_csv(r'C:\Users\admin\Desktop\2213557-MA 336\IRIS.csv')
```

## Exploring the Data Set

```
In [5]: # Display the first few rows of the dataset
print(IRIS_data.head())

# Check for missing values
print(IRIS_data.isnull().sum())

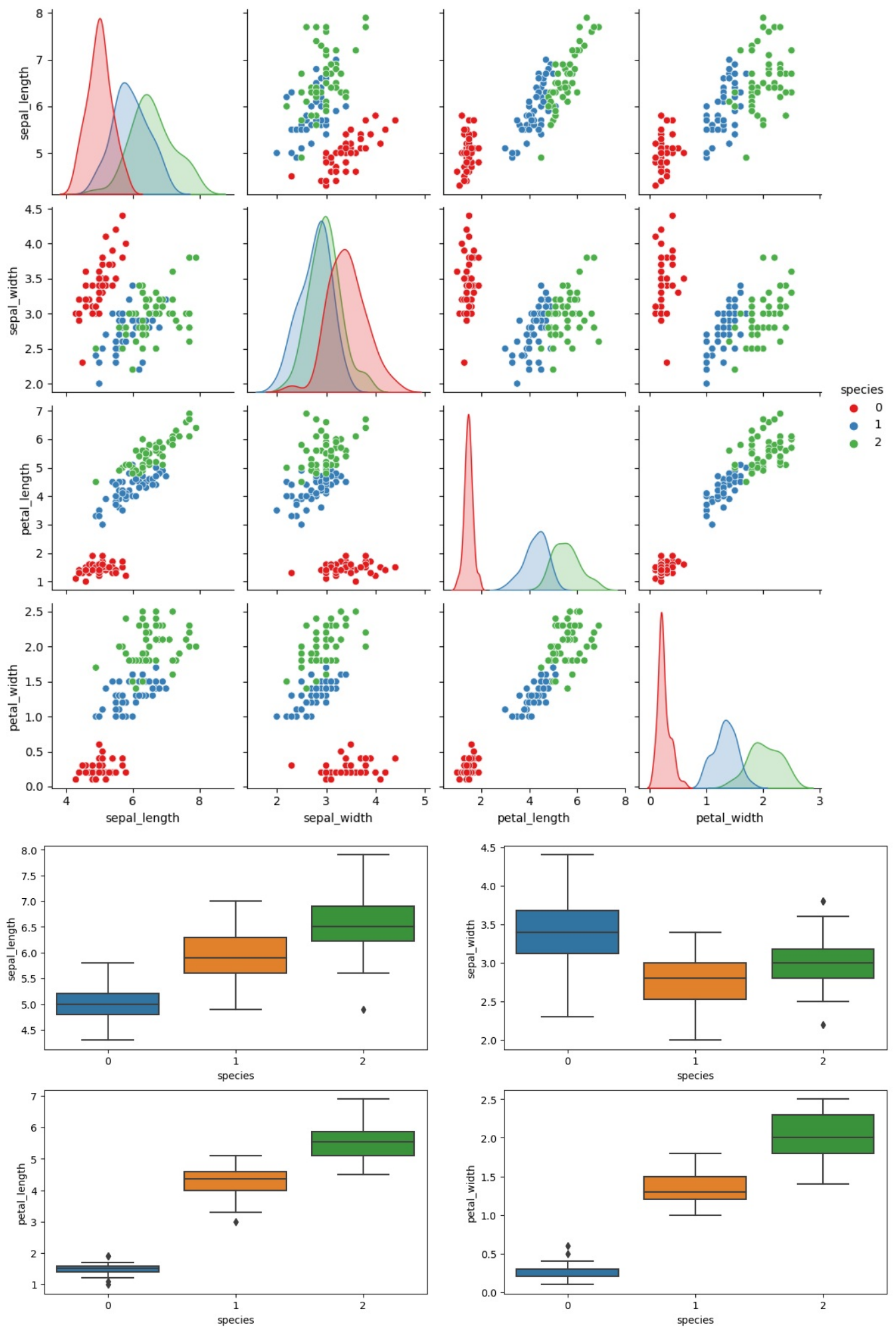
# Explore the distribution of classes
print(IRIS_data['species'].value_counts())
```

	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

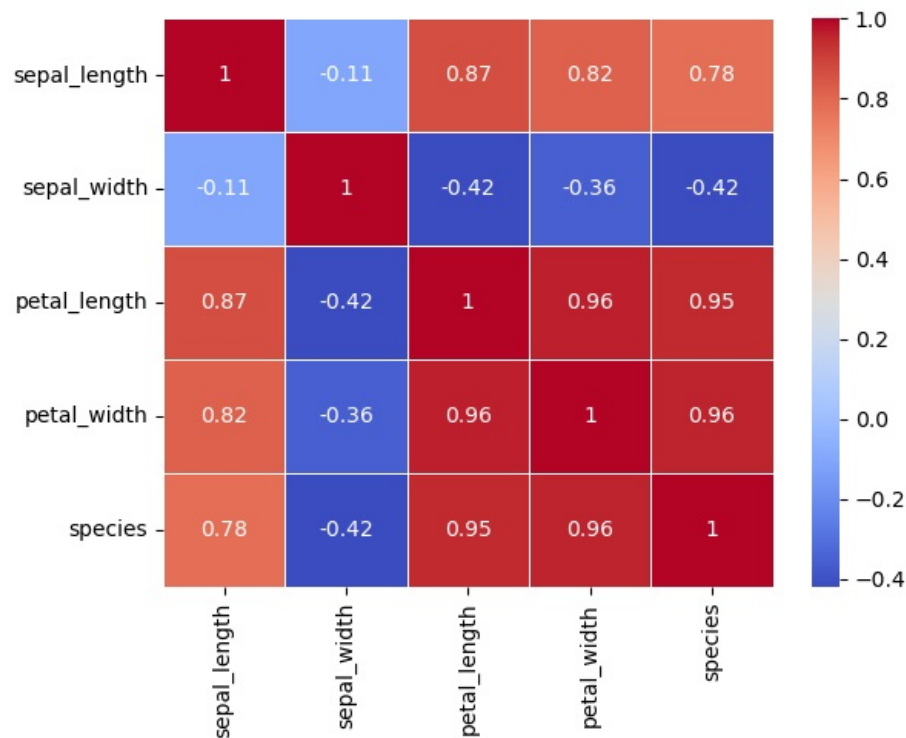
```
sepal_length    0
sepal_width     0
petal_length    0
petal_width     0
species         0
dtype: int64
Iris-setosa      50
Iris-versicolor  50
Iris-virginica   50
Name: species, dtype: int64
```

```
In [14]: # Pairplot to visualize relationships between features
sns.pairplot(IRIS_data, hue='species', markers='o', palette='Set1')
plt.show()

# Boxplot for each feature by species
plt.figure(figsize=(15, 8))
plt.subplot(2, 2, 1)
sns.boxplot(x='species', y='sepal_length', data=IRIS_data)
plt.subplot(2, 2, 2)
sns.boxplot(x='species', y='sepal_width', data=IRIS_data)
plt.subplot(2, 2, 3)
sns.boxplot(x='species', y='petal_length', data=IRIS_data)
plt.subplot(2, 2, 4)
sns.boxplot(x='species', y='petal_width', data=IRIS_data)
plt.show()
```



```
In [15]: # Correlation matrix to check for feature relationships
correlation_matrix = IRIS_data.corr()
sns.heatmap(correlation_matrix, annot=True, cmap='coolwarm', linewidths=.5)
plt.show()
```



## Preprocessing the Data Set

```
In [6]: # Encode the target variable
le = LabelEncoder()
IRIS_data['species'] = le.fit_transform(IRIS_data['species'])

# Split the data into features (X) and target (y)
X = IRIS_data.drop('species', axis=1)
y = IRIS_data['species']
```

Encoding the categorical target variable ('species') into numerical values and split the dataset into features (X) and target (y)

## Split the Dataset:

Splitting the dataset into training and testing sets.

```
In [7]: X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
```

## Train a Machine Learning Model:

Using Random Forest Classifier, to train on the training data.

```
In [8]: model = RandomForestClassifier(random_state=42)
model.fit(X_train, y_train)
```

```
Out[8]: RandomForestClassifier(random_state=42)
```

## Making Predictions

```
In [9]: y_pred = model.predict(X_test)
```

## Evaluating the Model

```
In [10]: accuracy = accuracy_score(y_test, y_pred)
print(f"Accuracy: {accuracy:.2f}")

classification_rep = classification_report(y_test, y_pred)
print("Classification Report:\n", classification_rep)
```

Accuracy: 1.00

Classification Report:

	precision	recall	f1-score	support
0	1.00	1.00	1.00	10
1	1.00	1.00	1.00	9
2	1.00	1.00	1.00	11
accuracy			1.00	30
macro avg	1.00	1.00	1.00	30
weighted avg	1.00	1.00	1.00	30

Here,The reported accuracy of 1.00 (100%) signifies exceptional performance by the machine learning model on the test dataset, demonstrating precise classification of Iris flowers into their respective species. The precision, recall, and F1-score metrics, all with perfect scores of 1.00 for each class (species 0, 1, and 2), highlight the model's accuracy and reliability in both positive and negative predictions