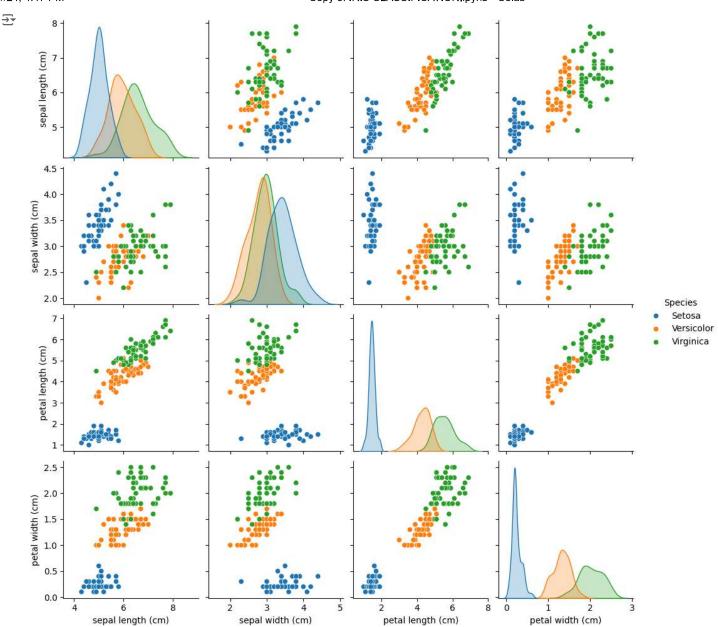
```
# Import libraries
import pandas as pd
import numpy as np
import seaborn as sns
import matplotlib.pyplot as plt
from sklearn.model_selection import train_test_split
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import accuracy_score, classification_report, confusion_matrix
from sklearn.datasets import load_iris
iris = load iris()
# Convert the dataset into a pandas DataFrame for easier analysis
iris_data = pd.DataFrame(iris.data, columns=iris.feature_names)
iris_data['Species'] = iris.target
# Replace numeric species labels with the actual names
species_map = {0: 'Setosa', 1: 'Versicolor', 2: 'Virginica'}
iris_data['Species'] = iris_data['Species'].map(species_map)
print("First few rows of the dataset:")
print(iris_data.head())
First few rows of the dataset:
        sepal length (cm) sepal width (cm) petal length (cm) petal width (cm) \
     0
                     5.1
                                       3.5
                                                          1.4
                                                                            0.2
     1
                     4.9
                                       3.0
                                                          1.4
                                                                            0.2
                     4.7
                                                                            0.2
     2
                                       3.2
                                                         1.3
     3
                     4.6
                                       3.1
                                                          1.5
                                                                            0.2
     4
                     5.0
                                       3.6
                                                          1.4
                                                                            0.2
      Species
     0 Setosa
     1 Setosa
     2 Setosa
     3 Setosa
     4 Setosa
# EDA: Pairplot to visualize features
sns.pairplot(data, hue='Species')
plt.show()
```



```
print("\nClassification Report:")
print(classification_report(y_test, y_pred))
```



Accuracy of the model: 1.00

Classification Report:

| precision | recall               | f1-score  | support   |
|-----------|----------------------|---|---|
| 1.00      | 1.00                 | 1.00  | 10  |
| 1.00      | 1.00                 | 1.00  | 9   |
| 1.00      | 1.00                 | 1.00  | 11  |
|           |                      | 1.00  | 30  |
| 1.00      | 1.00                 | 1.00  | 30  |
| 1.00      | 1.00                 | 1.00  | 30  |
|           | 1.00<br>1.00<br>1.00 | precision recall  1.00 1.00 1.00 1.00 1.00 1.00 1.00 1.00 | recision recall f1-score  1.00 1.00 1.00 1.00 1.00 1.00 1.00 1.00 |

```
print("\nConfusion Matrix:")
cm = confusion_matrix(y_test, y_pred)
species_names = list(species_map.values())
sns.heatmap(cm, annot=True, fmt='d', cmap='Blues', xticklabels=species_names, yticklabels=species_names)
plt.xlabel('Predicted')
plt.ylabel('Actual')
plt.title('Confusion Matrix')
plt.show()
```



## Confusion Matrix:

