

```

import pandas as pd
# Load dataset
df = pd.read_csv("iris_dataset.csv")
print(df.head())
# Explore
print(df.describe())
print("Target classes:", df["species"].unique())

```

```

      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
2                    4.7                3.2                1.3                0.2
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

      sepal length (cm)  sepal width (cm)  petal length (cm)  \
count          150.000000          150.000000          150.000000
mean             5.843333             3.057333             3.758000
std              0.828066             0.435866             1.765298
min              4.300000             2.000000             1.000000
25%              5.100000             2.800000             1.600000
50%              5.800000             3.000000             4.350000
75%              6.400000             3.300000             5.100000
max              7.900000             4.400000             6.900000

      petal width (cm)
count          150.000000
mean             1.199333
std              0.762238
min              0.100000
25%              0.300000
50%              1.300000
75%              1.800000
max              2.500000
Target classes: ['setosa' 'versicolor' 'virginica']

```

```
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.model_selection import train_test_split
from sklearn.linear_model import LogisticRegression
from sklearn.metrics import confusion_matrix, classification_report

df = pd.read_csv("iris_dataset.csv")

print("First 5 rows of the dataset:")
print(df.head())

sns.pairplot(df, hue="species", diag_kind="kde")
plt.suptitle("Iris Dataset Feature Relationships", y=1.02)
plt.show()

X = df.drop(columns=["species"])
y = df["species"]

X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2,
random_state=42)

model = LogisticRegression(max_iter=200)
model.fit(X_train, y_train)

y_pred = model.predict(X_test)

cm = confusion_matrix(y_test, y_pred)

print("\nConfusion Matrix:\n", cm)
print("\nClassification Report:\n", classification_report(y_test, y_pred))

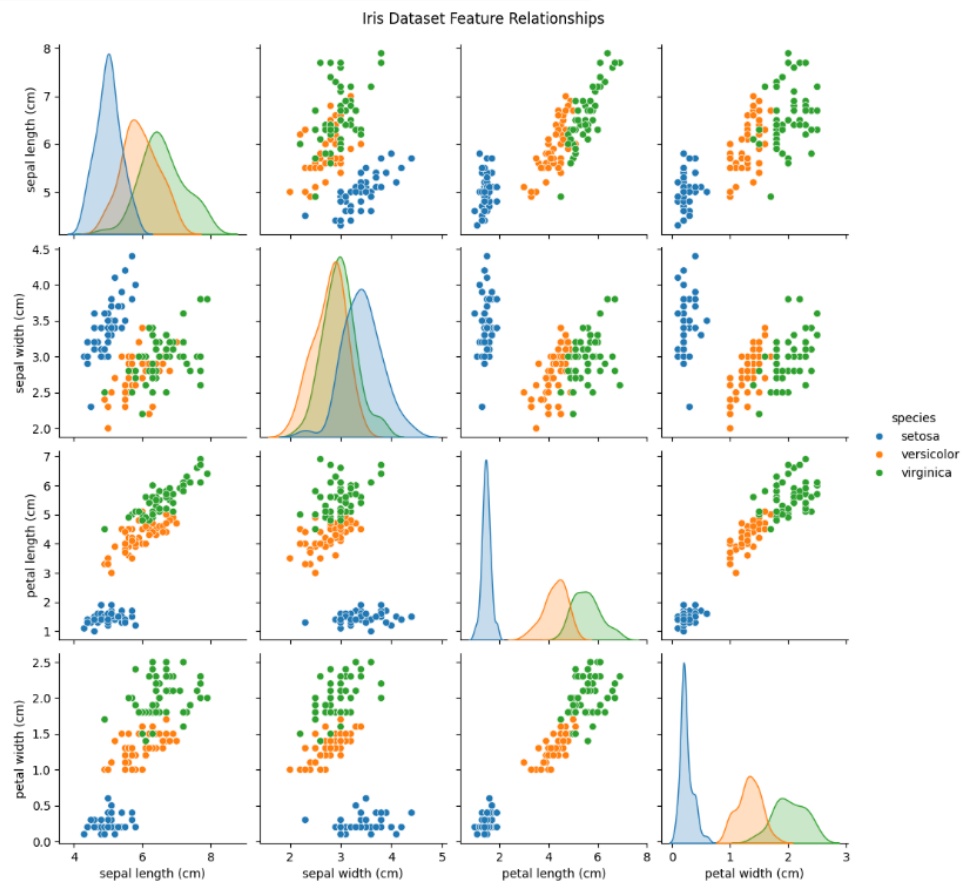
plt.figure(figsize=(6, 4))
sns.heatmap(cm, annot=True, cmap="Blues", fmt="d",
            xticklabels=model.classes_,
            yticklabels=model.classes_)
```

```
plt.xlabel("Predicted")
plt.ylabel("Actual")
plt.title("Confusion Matrix Heatmap")
plt.show()
```

```
plt.figure(figsize=(6, 4))
plt.scatter(range(len(y_test)), y_test, color='blue', label='Actual', alpha=0.7)
plt.scatter(range(len(y_pred)), y_pred, color='red', label='Predicted', alpha=0.5)
plt.title("Predicted vs Actual Species")
plt.xlabel("Sample Index")
plt.ylabel("Species")
plt.legend()
plt.show()
```

```
First 5 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
2                4.7                3.2                1.3                0.2
3                4.6                3.1                1.5                0.2
4                5.0                3.6                1.4                0.2

  species
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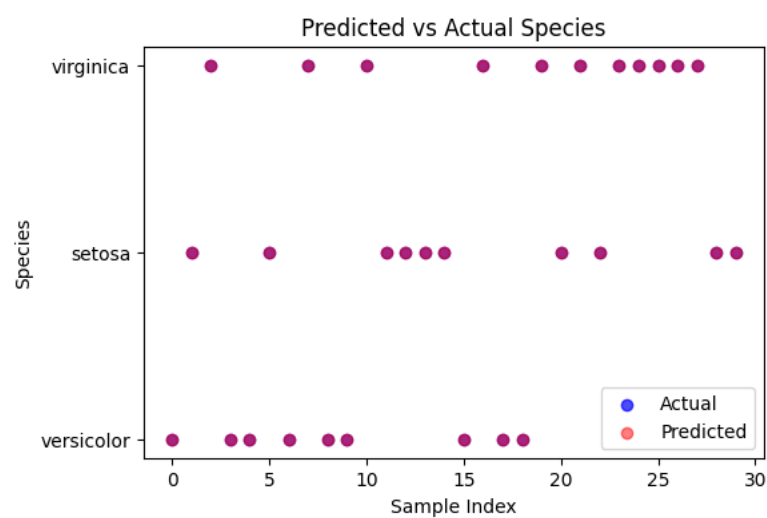
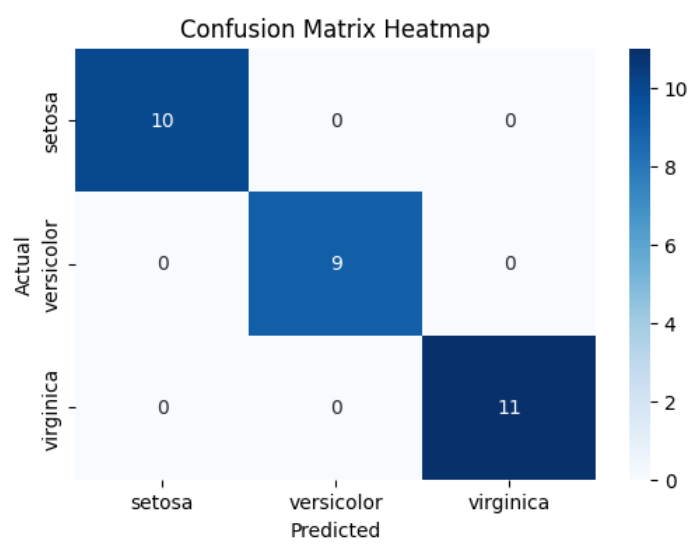


```
Confusion Matrix:
[[10  0  0]
 [ 0  9  0]
 [ 0  0 11]]

Classification Report:
              precision    recall  f1-score   support

   setosa      1.00      1.00      1.00        10
  versicolor  1.00      1.00      1.00         9
   virginica  1.00      1.00      1.00        11

 accuracy      1.00      1.00      1.00        30
  macro avg    1.00      1.00      1.00        30
 weighted avg  1.00      1.00      1.00        30
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