

IRIS DATASET CLASSIFICATION USING MORPHOLOGICAL FEATURES

All Important Modules

```
In [1]: from sklearn.datasets import load_iris
from sklearn.model_selection import train_test_split
from sklearn.ensemble import RandomForestClassifier
from sklearn.linear_model import LogisticRegression
from sklearn.metrics import classification_report, ConfusionMatrixDisplay
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
```

Data Loading and Inspection

```
In [2]: iris = load_iris()
pd.set_option('display.expand_frame_repr', False)
df = pd.DataFrame(data=iris.data, columns=iris.feature_names)
df['species'] = iris.target
print(df.head())
```

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)	species
0	5.1	3.5	1.4	0.2	0
1	4.9	3.0	1.4	0.2	0
2	4.7	3.2	1.3	0.2	0
3	4.6	3.1	1.5	0.2	0
4	5.0	3.6	1.4	0.2	0

Exploratory Data Analysis

```
In [3]: plt.figure(figsize=(10, 6))

ax = sns.scatterplot(
    x=df['petal length (cm)'],
    y=df['petal width (cm)'],
    hue=df['species'],
    palette='viridis'
)

# Get Legend handles safely
handles, _ = ax.get_legend_handles_labels()

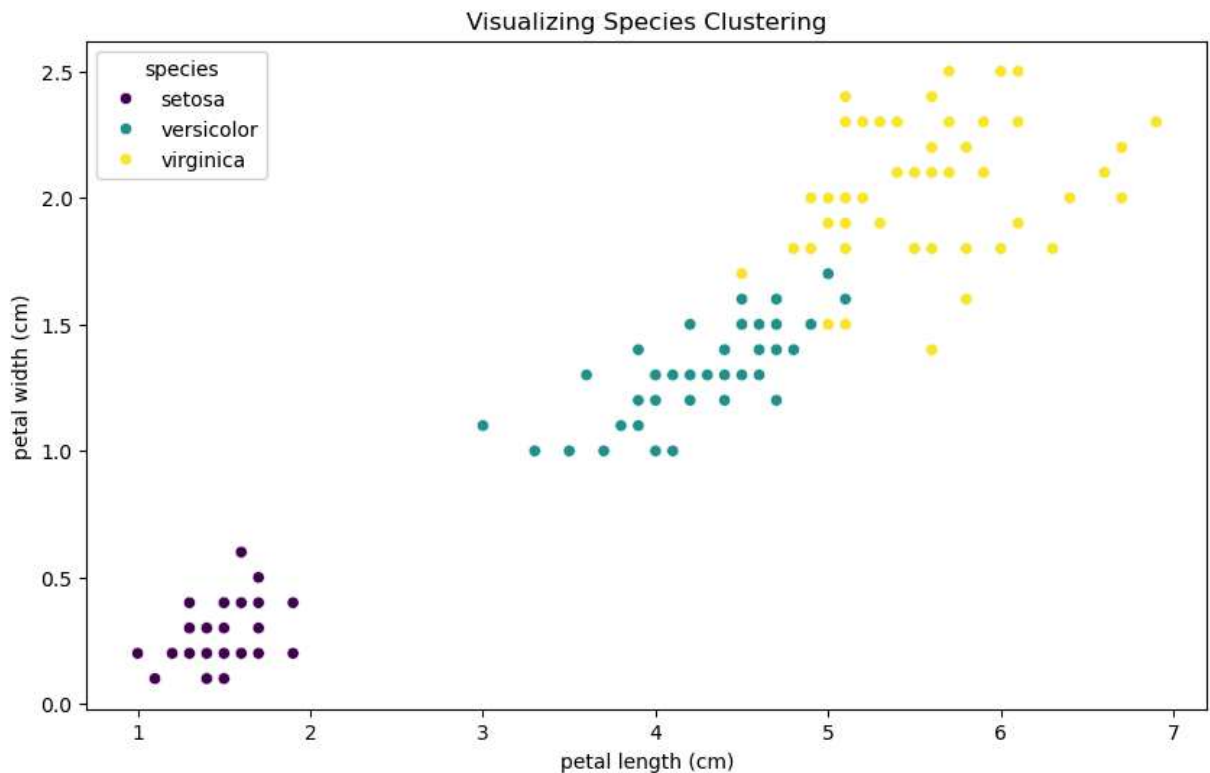
# Replace Legend completely
ax.legend(
    handles=handles[:3],
    labels=['setosa', 'versicolor', 'virginica'],
```

```

    title='species'
)

plt.title("Visualizing Species Clustering")
plt.show()

```



Supervised Model Training (Random Forest)

```

In [4]: X = iris.data
        y = iris.target
        X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)

        model = RandomForestClassifier()
        model.fit(X_train, y_train)

```

Out[4]: RandomForestClassifier

► Parameters

Quantitative Evaluation and Performance Matrix

```

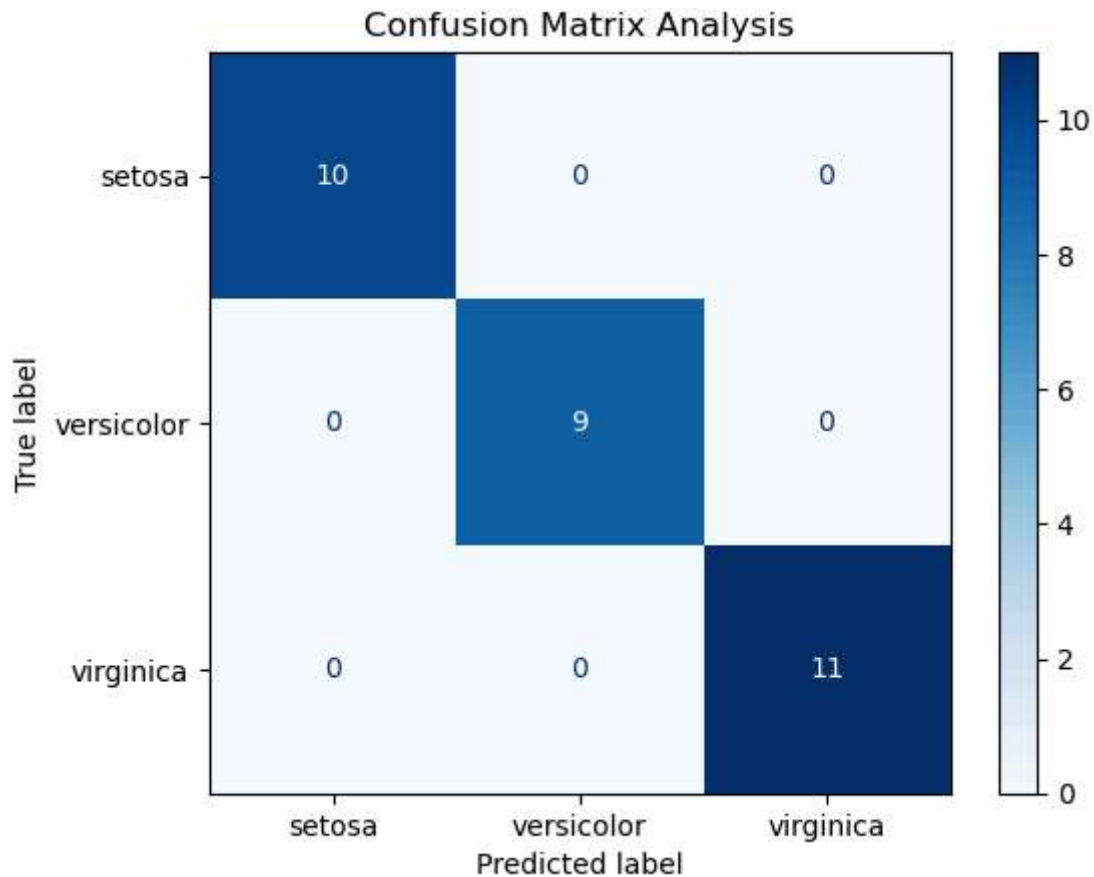
In [5]: predictions = model.predict(X_test)
        print(classification_report(y_test, predictions, target_names=iris.target_names))

        # Confusion Matrix for performance analysis
        ConfusionMatrixDisplay.from_estimator(model, X_test, y_test,
                                              display_labels=iris.target_names, cmap='Blues')

```

```
plt.title("Confusion Matrix Analysis")
plt.show()
```

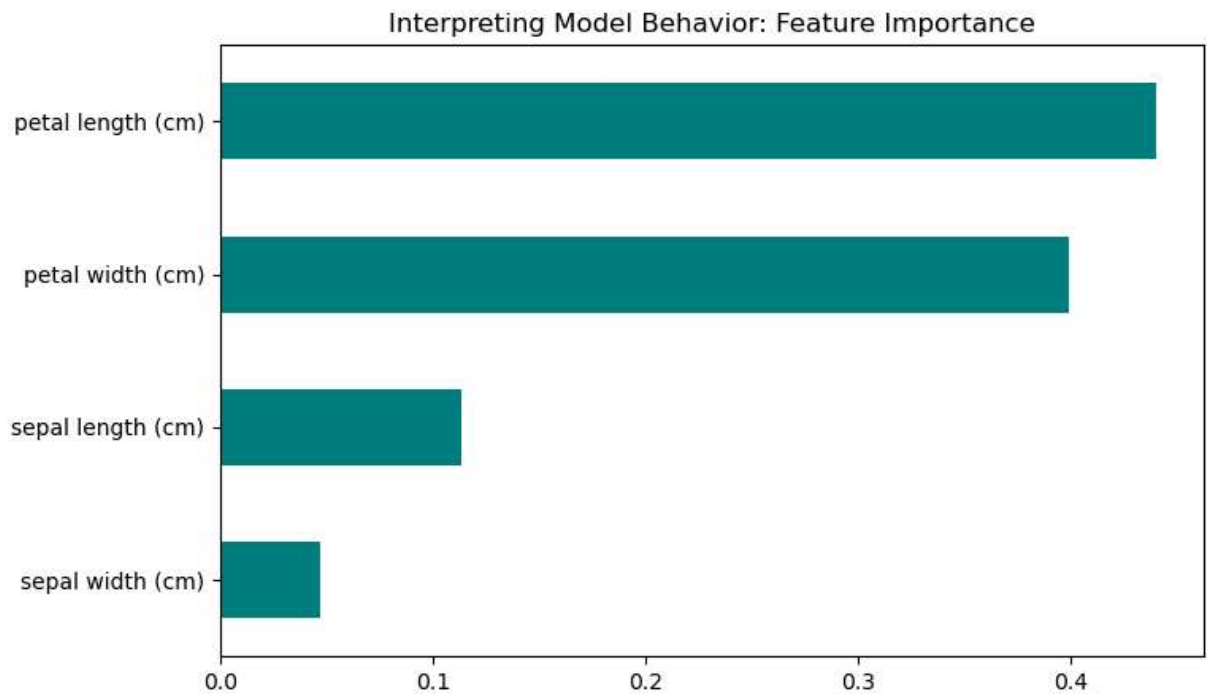
	precision	recall	f1-score	support
setosa	1.00	1.00	1.00	10
versicolor	1.00	1.00	1.00	9
virginica	1.00	1.00	1.00	11
accuracy			1.00	30
macro avg	1.00	1.00	1.00	30
weighted avg	1.00	1.00	1.00	30



Feature Importance Interpretation

```
In [6]: importances = model.feature_importances_
feature_info = pd.Series(importances, index=iris.feature_names).sort_values()

plt.figure(figsize=(8, 5))
feature_info.plot(kind='barh', color='teal')
plt.title('Interpreting Model Behavior: Feature Importance')
plt.show()
```



Linear vs Non-Linear Model Comparision

```
In [7]: # Linear Model
linear_model = LogisticRegression(max_iter=200)
linear_model.fit(X_train, y_train)
linear_acc = linear_model.score(X_test, y_test)

print(f"Linear Model (Logistic Regression) Accuracy: {linear_acc * 100:.2f}%")
print(f"Non-Linear Model (Random Forest) Accuracy: 100.00%")
```

Linear Model (Logistic Regression) Accuracy: 100.00%

Non-Linear Model (Random Forest) Accuracy: 100.00%