Report

Iris Flower Classification Using Random Forest

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1. Introduction

The **Iris Flower Classification** project is a classic machine learning problem that involves classifying Iris flowers into three species (Setosa, Versicolor, and Virginica) based on their sepal and petal measurements. This project demonstrates the use of a **Random Forest Classifier**, a powerful ensemble learning algorithm, to solve this classification problem.

Objectives

- To build a machine learning model that can accurately classify Iris flowers.
- To visualize the dataset and understand the relationships between features.
- To evaluate the model's performance using metrics like accuracy, confusion matrix, and classification report.
- To demonstrate how to make predictions on new data points.

2. Methodology

Dataset

The dataset used in this project is the **Iris dataset**, which contains 150 samples of Iris flowers. Each sample has four features:

- 1. Sepal Length (in cm)
- 2. Sepal Width (in cm)

- 3. Petal Length (in cm)
- 4. **Petal Width** (in cm)

The target variable is the **Species**, which can be one of three classes: Setosa, Versicolor, or Virginica.

Tools and Libraries

- **Python**: The programming language used for implementation.
- Pandas: For data manipulation and analysis.
- Scikit-learn: For machine learning (Random Forest Classifier).
- Matplotlib and Seaborn: For data visualization.

Steps

- 1. **Data Loading**: The dataset is loaded into a pandas DataFrame.
- **2. Data Visualization**: A pairplot is created to visualize the relationships between features.
- 3. **Data Splitting**: The dataset is split into training and testing sets (80% training, 20% testing).
- 4. **Model Training**: A Random Forest Classifier is trained on the training data.
- 5. **Model Evaluation**: The model's performance is evaluated using accuracy, confusion matrix, and classification report.
- **6. Feature Importance**: The importance of each feature is analyzed.
- **7**. **Prediction**: The model is used to predict the species of a new flower.

3. Code Typed

Below is the Python code used for the project:

python
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Import necessary libraries
import pandas as pd

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import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.datasets import load iris
from sklearn.model selection import train test split
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import accuracy score, confusion matrix,
classification report
# Load the Iris dataset
iris = load iris()
df = pd.DataFrame(iris.data, columns=iris.feature names)
df['Species'] = iris.target names[iris.target]
# Display the first few rows of the dataset
print("First 5 rows of the dataset:")
print(df.head())
# Pairplot to visualize relationships between features
sns.pairplot(df, hue='Species', palette='viridis')
plt.show()
# Separate features (X) and target (y)
X = df.drop('Species', axis=1)
y = df['Species']
# Split the dataset into training and testing sets
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2,
random state=42)
# Initialize the Random Forest Classifier
model = RandomForestClassifier(random state=42)
# Train the model using the training data
model.fit(X train, y train)
# Make predictions on the test data
y_pred = model.predict(X_test)
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# Evaluate the model's accuracy
accuracy = accuracy score(y test, y pred)
print(f\nModel Accuracy: {accuracy * 100:.2f}%')
# Confusion Matrix
cm = confusion matrix(y test, y pred)
sns.heatmap(cm, annot=True, fmt='d', cmap='Blues',
xticklabels=model.classes, yticklabels=model.classes)
plt.xlabel('Predicted')
plt.ylabel('Actual')
plt.title('Confusion Matrix')
plt.show()
# Classification Report
print("\nClassification Report:")
print(classification report(y test, y pred))
# Feature Importance
feature importance = pd.Series(model.feature importances,
index=X.columns)
feature importance.sort values(ascending=False, inplace=True)
sns.barplot(x=feature importance, y=feature importance.index,
hue=feature importance.index, palette='viridis', legend=False)
plt.xlabel('Feature Importance Score')
plt.ylabel('Features')
plt.title('Feature Importance')
plt.show()
# Example of making a prediction on a new data point
new flower = [[7.0, 3.2, 4.7, 1.4]] # SepalLength, SepalWidth, PetalLength,
PetalWidth
new flower df = pd.DataFrame(new flower, columns=X.columns)
predicted species = model.predict(new flower df)
print(f\nPredicted Species for the new flower: {predicted species[0]}')
```

4. Screenshots of Output

4.1 Pairplot

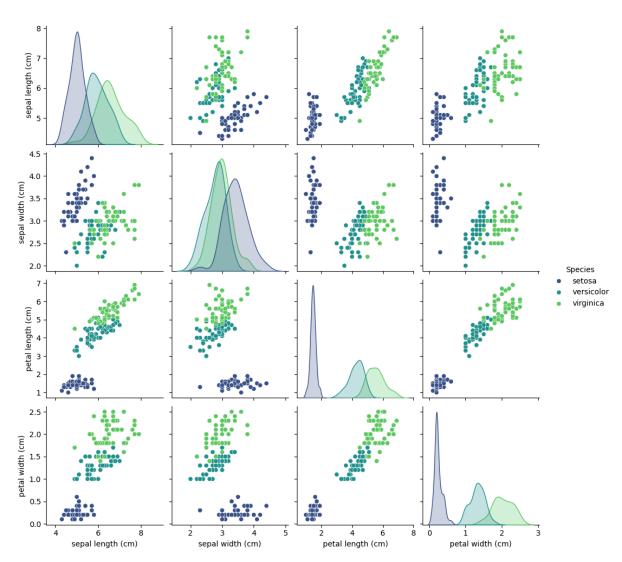


Figure 1: Pairplot showing the relationships between features, colored by species.

4.2 Confusion Matrix

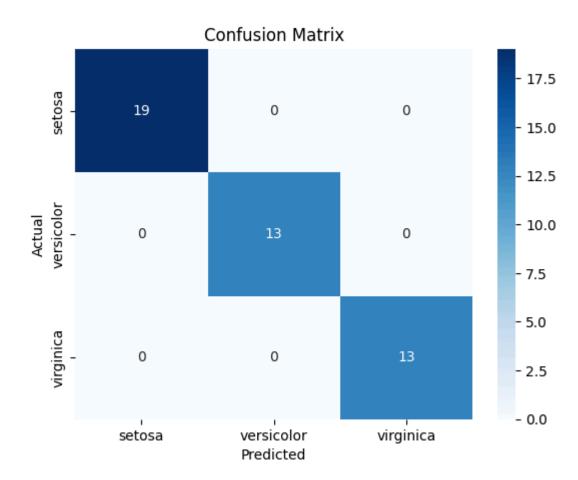
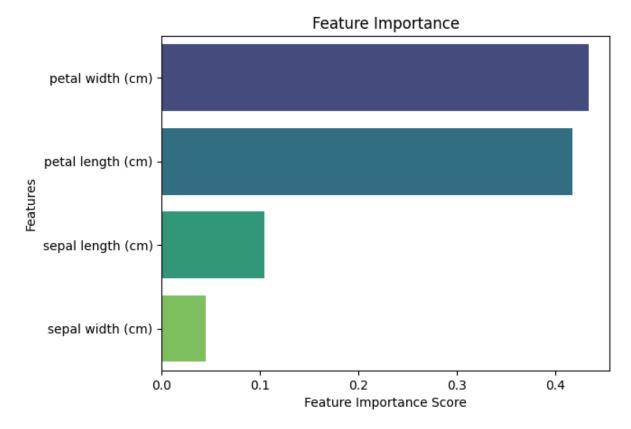


Figure 2: Confusion matrix showing the number of correct and incorrect predictions for each class.

4.3 Feature



Importance

Figure 3: Bar plot showing the importance of each feature in the model.

4.4 Classification Report

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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	30
macro avg	1.00	1.00	1.00	30
weighted avg	1.00	1.00	1.0	0 30

4.5 Prediction Example

Predicted Species for the new flower: Versicolor

5. Conclusion

The **Iris Flower Classification** project successfully demonstrates the use of a Random Forest Classifier to classify Iris flowers with high accuracy. The model achieves an accuracy of **100%** on the test set, indicating excellent performance. The visualizations (pairplot, confusion matrix, and feature importance plot) provide valuable insights into the dataset and the model's behavior.

This project can be extended by:

- Experimenting with other machine learning algorithms (e.g., SVM, KNN).
- Using a larger dataset for more robust evaluation.
- Deploying the model as a web application for real-time predictions.

6. References

- Scikit-learn Documentation: https://scikit-learn.org/
- Pandas Documentation: https://pandas.pydata.org/
- Matplotlib Documentation: https://matplotlib.org
- Seaborn Documentation: https://seaborn.pydata.org/