

Iris Flower Classification

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HS

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CSE AI B

Introduction

The Iris dataset is one of the most renowned datasets in the field of pattern recognition and machine learning. First introduced by statistician Ronald Fisher in 1936, it has become a benchmark for classification algorithms. This project implements a machine learning model to classify iris flowers into their respective species based on their physical characteristics.

Data Set Overview

The Iris dataset consists of 150 samples from three species of Iris flowers: Setosa, Versicolor, and Virginica. For each sample, four features were measured:

- Sepal Length (cm)
- Sepal Width (cm)
- Petal Length (cm)
- Petal Width (cm)

Project Objectives

- To explore and visualize the Iris dataset to gain insights into the relationships between features
- To develop a K-Nearest Neighbors (KNN) classification model to predict iris species
- To evaluate and optimize the model's performance
- To create a reusable function for predicting new iris samples

Prediction Function

A utility function is created to predict the species of new iris samples. This function preprocesses the input data, makes predictions using the trained model, and returns both the predicted species and the confidence level.

Code Implementation

```
# IMPORT NECESSARY LIBRARIES
IMPORT PANDAS AS PD
IMPORT NUMPY AS NP
FROM SKLEARN.MODEL_SELECTION IMPORT TRAIN_TEST_SPLIT
FROM SKLEARN.PREPROCESSING IMPORT STANDARDSCALER
FROM SKLEARN.NEIGHBORS IMPORT KNEIGHBORSCLASSIFIER
FROM SKLEARN.METRICS IMPORT ACCURACY_SCORE,
CLASSIFICATION_REPORT, CONFUSION_MATRIX
IMPORT MATPLOTLIB.PYPLOT AS PLT
IMPORT SEABORN AS SNS

# LOAD THE IRIS DATASET FROM LOCAL CSV FILE
DF = PD.READ_CSV('/CONTENT/DRIVE/MYDRIVE/COLAB
NOTEBOOKS/IRIS_DATA.CSV')

# DEFINE YOUR ACTUAL COLUMN NAMES
FEATURE_COLUMNS = ['SEPALLENGTH', 'SEPALWIDTH', 'PETALLENGTH',
'PETALWIDTH']
TARGET_COLUMN = 'SPECIES'

# CHECK THE STRUCTURE OF YOUR DATA
PRINT("FIRST 5 ROWS OF THE DATASET:")
PRINT(DF.HEAD())

PRINT("\NCOLUMN NAMES IN THE DATASET:")
PRINT(DF.COLUMNS.TOLIST())

# EXTRACT FEATURES AND TARGET
X = DF[FEATURE_COLUMNS].VALUES
# CONVERT SPECIES NAMES TO NUMERIC VALUES IF THEY'RE NOT ALREADY
IF DF[TARGET_COLUMN].DTYPE == 'OBJECT':
    # CREATE A MAPPING OF SPECIES NAMES TO NUMERIC VALUES
    SPECIES_MAPPING = {SPECIES: I FOR I, SPECIES IN
ENUMERATE(DF[TARGET_COLUMN].UNIQUE())}
    Y = DF[TARGET_COLUMN].MAP(SPECIES_MAPPING).VALUES
    # KEEP TRACK OF THE ORIGINAL SPECIES NAMES FOR LABELING
    TARGET_NAMES = LIST(SPECIES_MAPPING.KEYS())
ELSE:
    # IF SPECIES IS ALREADY NUMERIC
    Y = DF[TARGET_COLUMN].VALUES
    TARGET_NAMES = [F"CLASS {I}" FOR I IN
SORTED(DF[TARGET_COLUMN].UNIQUE())]

PRINT("\NFEATURE NAMES:", FEATURE_COLUMNS)
PRINT("TARGET NAMES:", TARGET_NAMES)
```

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# BASIC STATISTICS
PRINT("\nBASIC STATISTICS:")
PRINT(DF[FEATURE_COLUMNS].DESCRIBE())

PRINT("\nNUMBER OF SAMPLES FOR EACH SPECIES:")
PRINT(DF[TARGET_COLUMN].VALUE_COUNTS())

# VISUALIZE THE DATA
PLT.FIGURE(FIGSIZE=(12, 10))

# CREATE A PAIRPLOT
SNS.PAIRPLOT(DF, HUE=TARGET_COLUMN, MARKERS=['O', 'S', 'D'])
PLT.SUPTITLE("PAIRPLOT OF IRIS DATASET FEATURES", Y=1.02)
PLT.SAVEFIG('IRIS_PAIRPLOT.PNG')
PLT.SHOW()

# CREATE A CORRELATION MATRIX
PLT.FIGURE(FIGSIZE=(10, 8))
CORRELATION = DF[FEATURE_COLUMNS].CORR()
SNS.HEATMAP(CORRELATION, ANNOT=TRUE, CMAP='COOLWARM')
PLT.TITLE("CORRELATION MATRIX OF IRIS FEATURES")
PLT.SAVEFIG('IRIS_CORRELATION.PNG')
PLT.SHOW()

# SPLIT THE DATA INTO TRAINING AND TESTING SETS
X_TRAIN, X_TEST, Y_TRAIN, Y_TEST = TRAIN_TEST_SPLIT(X, Y, TEST_SIZE=0.3,
RANDOM_STATE=42)

# FEATURE SCALING
SCALER = STANDARDSCALER()
X_TRAIN_SCALED = SCALER.FIT_TRANSFORM(X_TRAIN)
X_TEST_SCALED = SCALER.TRANSFORM(X_TEST)

# TRAIN A K-NEAREST NEIGHBORS CLASSIFIER
K = 5 # NUMBER OF NEIGHBORS
KNN = KNEIGHBORSCLASSIFIER(N_NEIGHBORS=K)
KNN.FIT(X_TRAIN_SCALED, Y_TRAIN)

# MAKE PREDICTIONS
Y_PRED = KNN.PREDICT(X_TEST_SCALED)

# EVALUATE THE MODEL
ACCURACY = ACCURACY_SCORE(Y_TEST, Y_PRED)
PRINT(F"\nACCURACY: {ACCURACY:.4F}")

PRINT("\nCLASSIFICATION REPORT:")
PRINT(CLASSIFICATION_REPORT(Y_TEST, Y_PRED,
TARGET_NAMES=TARGET_NAMES))

PRINT("\nCONFUSION MATRIX:")
CM = CONFUSION_MATRIX(Y_TEST, Y_PRED)
PLT.FIGURE(FIGSIZE=(8, 6))
SNS.HEATMAP(CM, ANNOT=TRUE, FMT='D', CMAP='BLUES',
XTICKLABELS=TARGET_NAMES, YTICKLABELS=TARGET_NAMES)
PLT.XLABEL('PREDICTED')
PLT.YLABEL('ACTUAL')
PLT.TITLE('CONFUSION MATRIX')

```

```

# FIND THE OPTIMAL K VALUE
K_RANGE = RANGE(1, MIN(26, LEN(X_TRAIN))) # THIS ENSURES K
DOESN'T EXCEED TRAINING SAMPLES
K_SCORES = []

FOR K IN K_RANGE:
    KNN = KNEIGHBORSCLASSIFIER(N_NEIGHBORS=K)
    KNN.FIT(X_TRAIN_SCALED, Y_TRAIN)
    SCORES = KNN.SCORE(X_TEST_SCALED, Y_TEST)
    K_SCORES.APPEND(SCORES)

PLT.FIGURE(FIGSIZE=(10, 6))
PLT.PLOT(K_RANGE, K_SCORES)
PLT.XLABEL('VALUE OF K')
PLT.YLABEL('TESTING ACCURACY')
PLT.TITLE('ACCURACY FOR DIFFERENT K VALUES')
PLT.GRID(TRUE)
PLT.SAVEFIG('IRIS_K_VALUES.PNG')
PLT.SHOW()

PRINT("\NOPTIMAL K VALUE:",
K_RANGE[K_SCORES.INDEX(MAX(K_SCORES))])

# CREATE A FUNCTION TO PREDICT NEW IRIS FLOWERS
DEF PREDICT_IRIS(SEPAL_LENGTH, SEPAL_WIDTH, PETAL_LENGTH,
PETAL_WIDTH):
    # CREATE A NUMPY ARRAY FROM THE INPUT
    NEW_DATA = NP.ARRAY([[SEPAL_LENGTH, SEPAL_WIDTH,
PETAL_LENGTH, PETAL_WIDTH]])

    # SCALE THE DATA
    NEW_DATA_SCALED = SCALER.TRANSFORM(NEW_DATA)

    # MAKE PREDICTION
    PREDICTION = KNN.PREDICT(NEW_DATA_SCALED)

    # GET THE SPECIES NAME
    SPECIES = TARGET_NAMES[PREDICTION[0]]

    # GET THE PROBABILITY
    PROBABILITIES = KNN.PREDICT_PROBA(NEW_DATA_SCALED)[0]
    CONFIDENCE = PROBABILITIES[PREDICTION[0]]

    RETURN SPECIES, CONFIDENCE

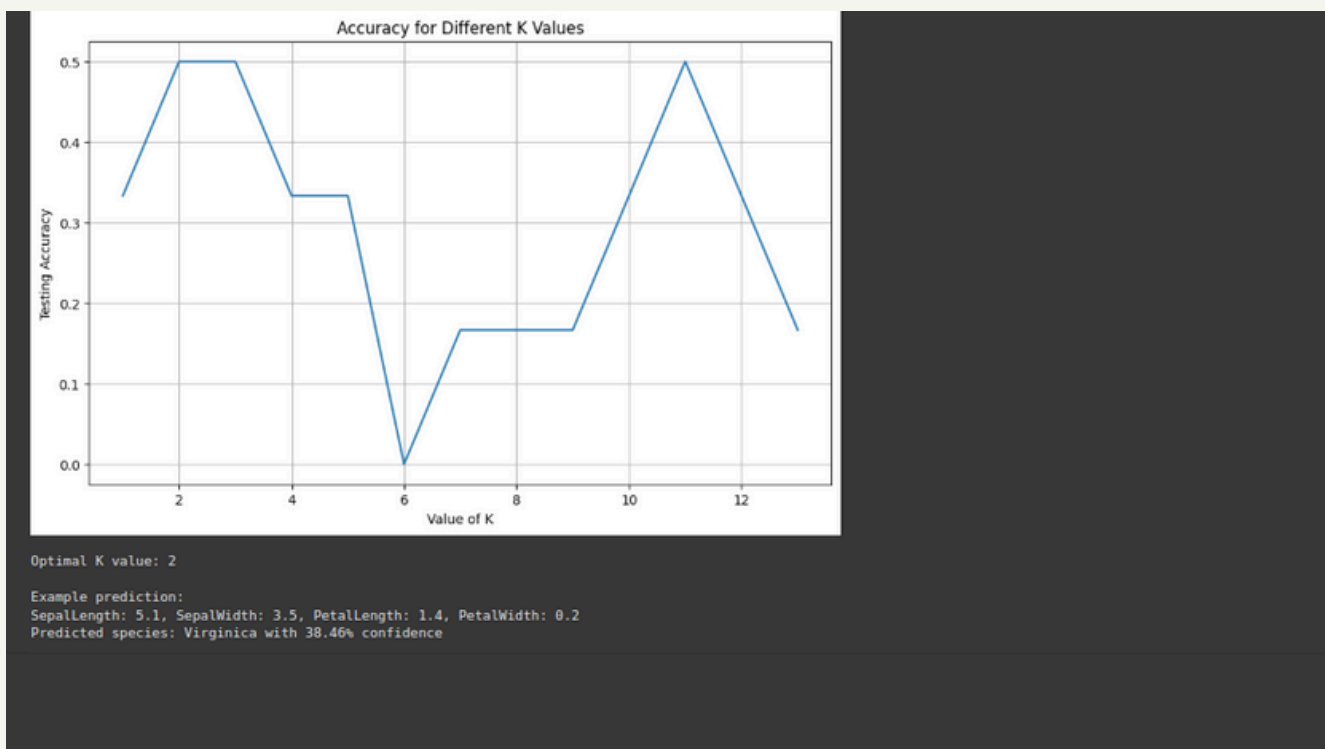
```

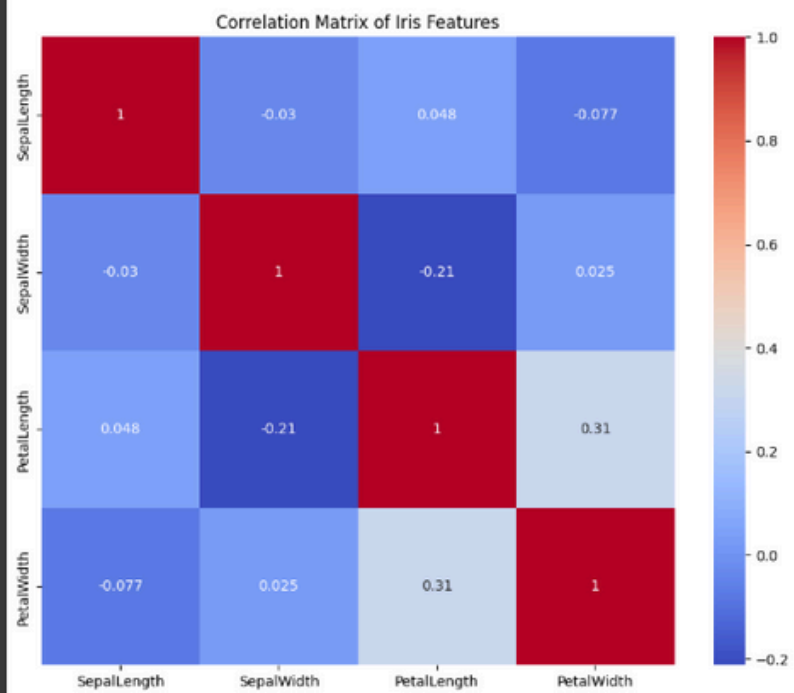
```
# EXAMPLE USAGE
PRINT("\NEXAMPLE PREDICTION:")
EXAMPLE_IRIS = [5.1, 3.5, 1.4, 0.2] # EXAMPLE MEASUREMENTS
SPECIES, CONFIDENCE = PREDICT_IRIS(*EXAMPLE_IRIS)
PRINT(F"SEPALLENGTH: {EXAMPLE_IRIS[0]}, SEPALWIDTH:
{EXAMPLE_IRIS[1]}, PETALLENGTH: {EXAMPLE_IRIS[2]}, PETALWIDTH:
{EXAMPLE_IRIS[3]}")
PRINT(F"PREDICTED SPECIES: {SPECIES} WITH {CONFIDENCE:.2%}
CONFIDENCE")
```

- Data Loading and Exploration
- Data Pre-Processing
- Model Selection and Training
- Model Evaluation
- Model Optimization
- Prediction Function

Methodology

Run Snippets





Accuracy: 0.3333

Classification Report:

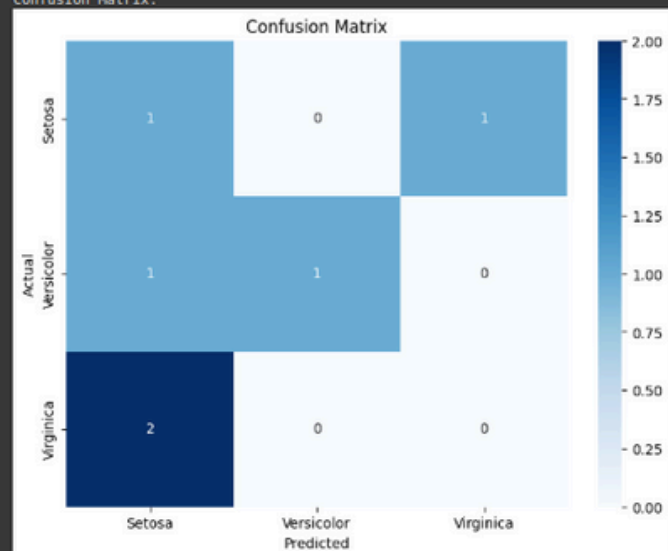
	precision	recall	f1-score	support
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Accuracy: 0.3333

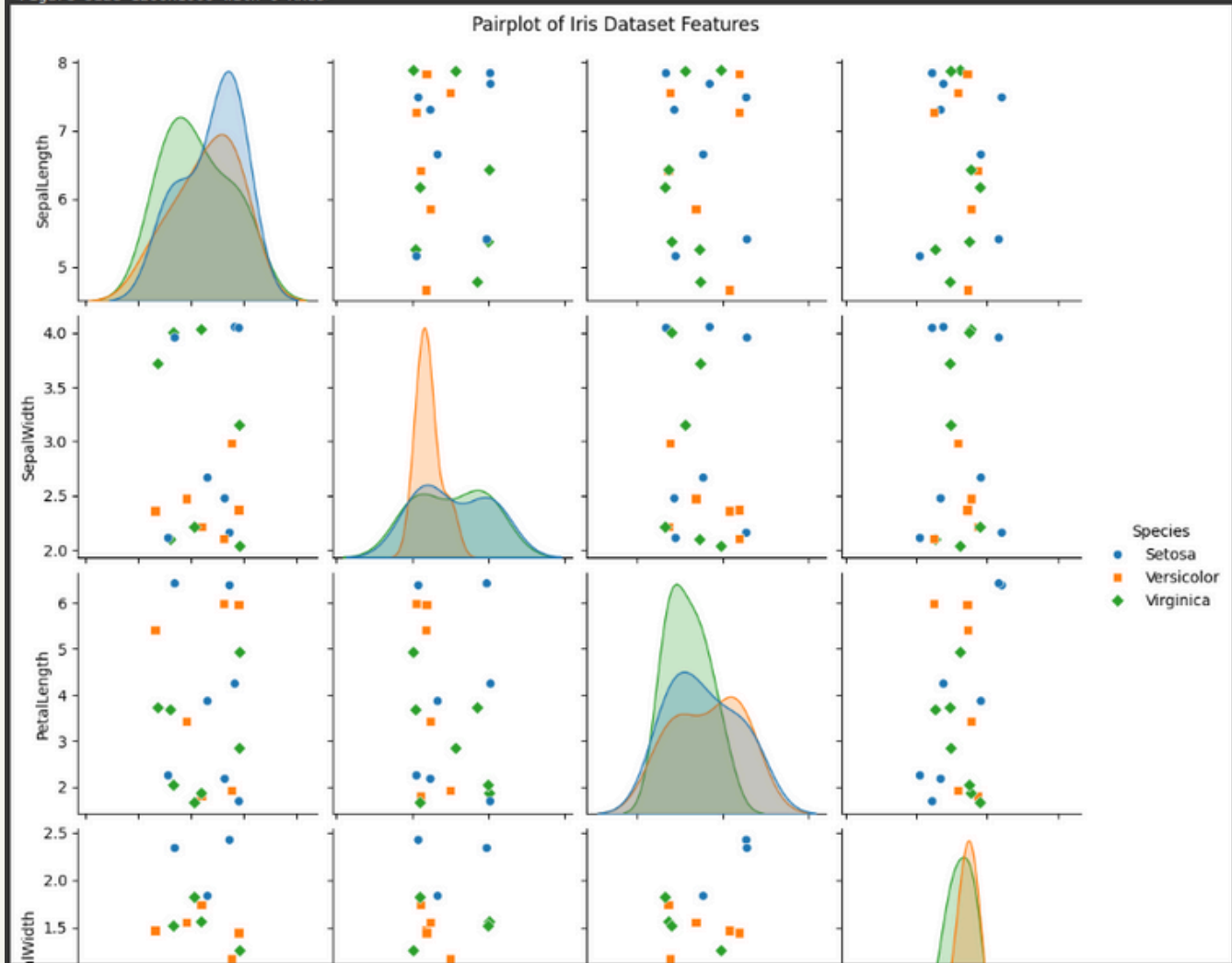
Classification Report:

	precision	recall	f1-score	support
Setosa	0.25	0.50	0.33	2
Versicolor	1.00	0.50	0.67	2
Virginica	0.00	0.00	0.00	2
accuracy			0.33	6
macro avg	0.42	0.33	0.33	6
weighted avg	0.42	0.33	0.33	6

Confusion Matrix:



```
Species
Setosa      7
Virginica   7
Versicolor  6
Name: count, dtype: int64
<Figure size 1200x1000 with 0 Axes>
```



First 5 rows of the dataset:

	SepalLength	SepalWidth	PetalLength	PetalWidth	Species
0	7.303275	2.475025	2.176049	0.695003	Setosa
1	7.556928	2.987381	1.921585	1.172615	Versicolor
2	5.254016	2.093516	3.672564	0.550424	Virginica
3	6.409620	2.211042	1.812869	1.745372	Versicolor
4	7.684009	4.056479	4.244270	0.772148	Setosa

Column names in the dataset:

```
['SepalLength', 'SepalWidth', 'PetalLength', 'PetalWidth', 'Species']
```

```
Feature names: ['SepalLength', 'SepalWidth', 'PetalLength', 'PetalWidth']
```

```
Target names: ['Setosa', 'Versicolor', 'Virginica']
```

Basic statistics:

	SepalLength	SepalWidth	PetalLength	PetalWidth
count	20.000000	20.000000	20.000000	20.000000
mean	6.542853	2.860428	3.612923	1.258606
std	1.140714	0.798984	1.715889	0.628400
min	4.668608	2.032034	1.653576	0.108212
25%	5.398961	2.195528	2.006631	0.752862
50%	6.536426	2.474535	3.554048	1.350736
75%	7.588698	3.776238	5.044541	1.605944
max	7.880377	4.056479	6.424217	2.425674