

# DEPARTMENT OF ARTIFICIAL INTELLIGENCE AND MACHINE LEARNING

CB19342 – COMPUTATIONAL STATISTICS

LAB RECORD - III YEAR

FIFTH SEMESTER2024-2025

**ODD SEMESTER** 

#### 1. PREDICTING HOUSE PRICES

EX.N0:1	Predicting House Prices
DATE: 24/07/2024	

**PROBLEM STATEMENT:** Build a regression model to predict house prices based on features like location, size, and amenities.

**PYTHON CONCEPTS:** Functions, classes, numeric types, sequences.

**<u>VISUALIZATION:</u>** Plotting regression line, residual plots.

**MULTIVARIATE ANALYSIS:** Multiple regression.

**DATASET:** Kaggle House Prices

#### **ALGORITHM:**

Step 1: Start the program.

Step 2: Import necessary libraries.

Step 3: Load the dataset.

Step 4: Encode categorical variable, define feature & testing set.

Step 5: Split the dataset into training & testing set, create trained model.

Step 6: Print equal metric & test the cell.

#### **PROGRAM:**

import pandas as pd

from sklearn.preprocessing import LabelEncoder

from sklearn.model\_selection import train\_test\_split

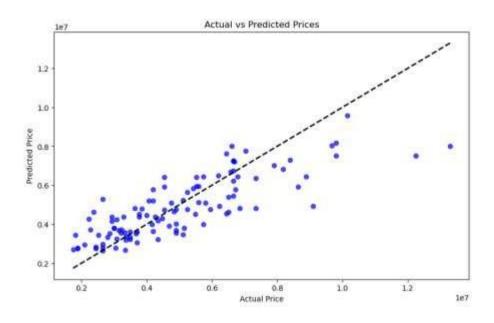
from sklearn.linear\_model import LinearRegression

```
from sklearn.metrics import r2_score, mean_absolute_error
import matplotlib.pyplot as plt
file_path =
'C:/Users/HEMA/Downloads/Housing.csv'
housing_data = pd.read_csv(file_path)
categorical_features = ['mainroad', 'guestroom', 'basement', 'hotwaterheating', 'airconditioning',
'prefarea', 'furnishingstatus']
le = LabelEncoder()
for feature in categorical_features:
housing_data[feature] = le.fit_transform(housing_data[feature])
X = housing_data.drop('price', axis=1)y = housing_data['price']
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
model = LinearRegression()
model.fit(X_train, y_train)
y_pred = model.predict(X_test)
r2 = r2\_score(y\_test, y\_pred)
mae = mean_absolute_error(y_test, y_pred)
plt.figure(figsize=(10, 6))
plt.scatter(y_test, y_pred, alpha=0.7, color='b')
plt.plot([y_test.min(), y_test.max()],
[y_test.min(), y_test.max()], 'k--', lw=2)
plt.xlabel('Actual Price')
plt.ylabel('Predicted Price')
plt.title('Actual vs Predicted Prices')
```

221501045

**COMPUTATIONAL STATISTICS** 

```
plt.show()
print(f'R-squared (R²): {r2}')
print(f'Mean Absolute Error (MAE): {mae}')
```



```
import numpy as np
test=np.array([ 7420,4,2,3,1,0,0,0,1,2,1,0]).reshape(-12,12)
model.predict(test)
```

array([8004072.41154001])

## **RESULT**:

Thus, the program for house price prediction is executed successfully.

#### 2. CUSTOMER SEGMENTATION FOR AN E-COMMERCE COMPANY

EX.N0: 2	Customer Segmentation for an E-commerce
DATE: 05/08/2024	Company

**PROBLEM STATEMENT:** Perform cluster analysis to segment customers based on purchasing behaviour.

**PYTHON CONCEPTS:** Data structures, file reading/writing.

**VISUALIZATION:** Cluster plots.

**MULTIVARIATE ANALYSIS:** Cluster analysis with k-means, hierarchical clustering.

**DATASET:** Online Retail Dataset

#### **ALGORITHM:**

Step 1: Start the program.

Step 2: Import necessary libraries.

Step 3: Load the dataset.

Step 4: Encode categorical variable, define feature & testing set.

Step 5: Split the dataset into training & testing set, create trained model.

Step 6: Print equal metric & test the cell.

#### **PROGRAM:**

import pandas as pd

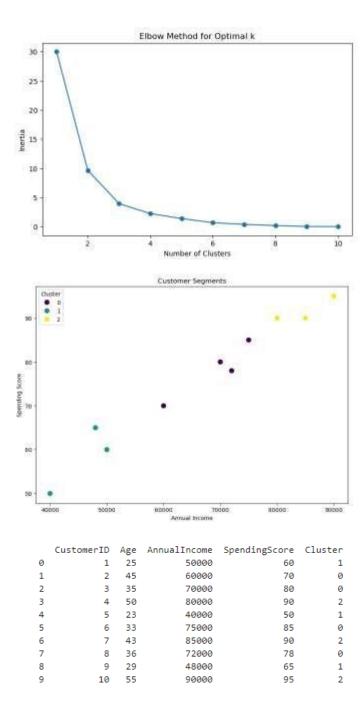
import numpy as np

from sklearn.preprocessing import StandardScaler

from sklearn.cluster import KMeans

import matplotlib.pyplot as plt

```
import seaborn as sns
import os
os.environ['OMP_NUM_THREADS'] = '1'
data = {'CustomerID': [1, 2, 3, 4, 5, 6, 7, 8, 9, 10],
'Age': [25, 45, 35, 50, 23, 33, 43, 36, 29, 55],
'AnnualIncome': [50000, 60000, 70000, 80000, 40000, 75000, 85000, 72000, 48000, 90000],
'SpendingScore': [60, 70, 80, 90, 50, 85, 90, 78, 65, 95] }
df = pd.DataFrame(data)
features = df[['Age', 'AnnualIncome', 'SpendingScore']]
scaler = StandardScaler()
scaled_features = scaler.fit_transform(features) inertia = []
k_range = range(1, 11) for k in k_range:
kmeans = KMeans(n_clusters=k, n_init=10, random_state=0)
kmeans.fit(scaled features)
inertia.append(kmeans.inertia_) plt.figure(figsize=(8, 5))
plt.plot(k_range, inertia, marker='o')
plt.xlabel('Number of Clusters') plt.ylabel('Inertia')
plt.title('Elbow Method for Optimal k') plt.show() optimal k = 3
kmeans = KMeans(n_clusters=optimal_k, n_init=10, random_state=0)
df['Cluster'] = kmeans.fit_predict(scaled_features)
plt.figure(figsize=(10, 7))
sns.scatterplot(data=df, x='AnnualIncome', y='SpendingScore', hue='Cluster', palette='viridis',
s=100)
plt.title('Customer Segments')
plt.xlabel('Annual Income')
plt.ylabel('Spending Score')
plt.legend(title='Cluster')
plt.show()
print(df)
```



## **RESULT**:

Thus, the program for Customer Segmentation for an E-commerce Company is executed successfully.

## 3. SENTIMENT ANALYSIS OF MOVIE REVIEWS

**EX.N0:3** 

## SENTIMENT ANALYSIS OF MOVIE REVIEWS

**DATE: 07/08/2024** 

**PROBLEM STATEMENT:** Classify movie reviews as positive or negative using text

Data.

**PYTHON CONCEPTS:** Text files, sequences, flow controls.

**VISUALIZATION:** Word cloud, bar plots.

**MULTIVARIATE ANALYSIS:** PCA for text data, logistic regression.

**DATASET:** IMDB Movie Reviews.

#### **ALGORITHM:**

Step 1: Start the program.

Step 2: Import necessary libraries.

Step 3: Load the dataset.

Step 4: Encode categorical variable, define feature & testing set.

Step 5: Split the dataset into training & testing set, create trained model.

Step 6: Print equal metric & test the cell.

#### **PROGRAM:**

import pandas as pd

import matplotlib.pyplot as plt

from wordcloud import WordCloud

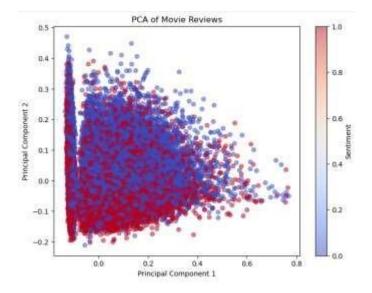
from sklearn.feature\_extraction.text import TfidfVectorizer

from sklearn.decomposition import PCA

from sklearn.linear\_model import LogisticRegression

```
from sklearn.metrics import classification_report, confusion_matrix
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import LabelEncoder
import nltk
from nltk.corpus import stopwords
from nltk.tokenize import word_tokenize
from nltk.stem import PorterStemmer
import seaborn as sns
nltk.download('punkt')
nltk.download('stopwords')
df = pd.read_csv('C:/Users/AASHI/Downloads/IMDB Dataset.csv')
stop_words = set(stopwords.words('english'))
stemmer = PorterStemmer()
def preprocess_text(text):
tokens = word_tokenize(text.lower())
tokens = [stemmer.stem(word) for word in tokens if word.isalpha() and word not in stop_words]
return ' '.join(tokens)
df['cleaned_review'] = df['review'].apply(preprocess_text)
vectorizer = TfidfVectorizer(max_features=5000)
X = vectorizer.fit_transform(df['cleaned_review']).toarray()
encoder = LabelEncoder()
y = encoder.fit_transform(df['sentiment'])
pca = PCA(n_components=2)
X_pca = pca.fit_transform(X)
plt.figure(figsize=(8, 6))
plt.scatter(X_pca[:, 0], X_pca[:, 1], c=y, cmap='coolwarm', alpha=0.5)
plt.title('PCA of Movie Reviews')
plt.xlabel('Principal Component 1')
plt.ylabel('Principal Component 2')
plt.colorbar(label='Sentiment')
plt.show()
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
model = LogisticRegression(max_iter=1000)
```

```
model.fit(X_train, y_train)
y_pred = model.predict(X_test)
print("Confusion Matrix:")
print(confusion_matrix(y_test, y_pred))
print("\nClassification Report:")
print(classification_report(y_test, y_pred))
positive_reviews = ' '.join(df[df['sentiment'] == 1]['cleaned_review'])
negative_reviews = ' '.join(df[df['sentiment'] == 0]['cleaned_review'])
plt.figure(figsize=(12, 6))
if len(positive_reviews.strip()) > 0:
plt.subplot(1, 2, 1)
plt.imshow(WordCloud(width=800, height=400,
background_color='white').generate(positive_reviews), interpolation='bilinear')
plt.title('Positive Reviews')
plt.axis('off')
else: print("No content available for positive reviews.")
if len(negative_reviews.strip()) > 0:
plt.subplot(1, 2, 2)
plt.imshow(WordCloud(width=800, height=400,
background_color='white').generate(negative_reviews), interpolation='bilinear')
plt.title('Negative Reviews')
plt.axis('off') else:
print("No content available for negative reviews.")
plt.show()
sns.countplot(x='sentiment', data=df)
plt.title('Sentiment Distribution')
plt.xlabel('Sentiment')
plt.ylabel('Count')
plt.show()
```



Confusion Matrix: [[4306 655] [ 511 4528]]

Classification Report:						
	precision recall		f1-score	support		
0	0.89	0.87	0.88	4961		
1	0.87	0.90	0.89	5039		
accuracy			0.88	10000		
macro avg	0.88	0.88	0.88	10000		
weighted avg	0.88	0.88	0.88	10000		

## **RESULT**:

Thus, the program for sentiment analysis of movie reviews is executed successfully.

#### 4. STOCK MARKET ANALYSIS

**EX.N0:4** 

## STOCK MARKET ANALYSIS

**DATE:** 14/08/2024

**PROBLEM STATEMENT:** Analyse stock market data to predict future stock prices.

**PYTHON CONCEPTS:** Data structures, file reading/writing, functions.

**<u>VISUALIZATION:</u>** Line plots, candlestick charts.

**MULTIVARIATE ANALYSIS:** Time series analysis, regression.

**DATASET:** Yahoo Finance Stock Data.

#### **ALGORITHM:**

Step 1: Start the program.

Step 2: Import necessary libraries.

Step 3: Load the dataset.

Step 4: Encode categorical variable, define feature & testing set.

Step 5: Split the dataset into training & testing set, create trained model.

Step 6: Print equal metric & test the cell.

#### **PROGRAM:**

import pandas as pd

import matplotlib.pyplot as plt

import mplfinance as mpf

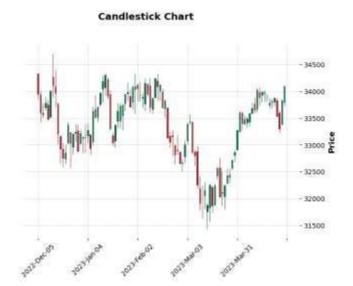
from statsmodels.tsa.arima.model import ARIMA

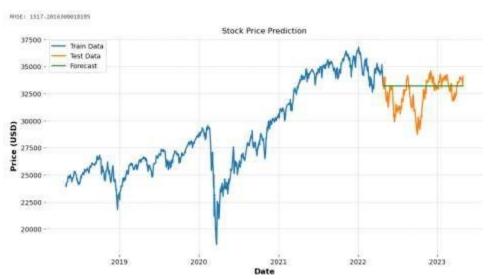
from sklearn.metrics import mean\_squared\_error

import numpy as np

file\_path = r'C:\Users\HEMA\Downloads\yahoo\_data.xlsx'

```
data = pd.read_excel(file_path, index_col='Date', parse_dates=True)
data.rename(columns={'Close*': 'Close', 'Adj Close**': 'Adj Close'}, inplace=True)
data.sort index(inplace=True)
data.ffill(inplace=True)
if 'Adj Close' in data.columns:
plt.figure(figsize=(12, 6))
plt.plot(data['Adj Close'], label='Adjusted Close Price')
plt.title('Adjusted Close Price Over Time')
plt.xlabel('Date')
plt.ylabel('Price (USD)')
plt.legend()
plt.show()
reduced_data = data[-100:] # Reduce data points for candlestick chart
mpf.plot(reduced_data, type='candle', style='charles', title='Candlestick Chart')
train_data, test_data = data['Adj Close'][:int(len(data)*0.8)], data['Adj Close'][int(len(data)*0.8):]
model = ARIMA(train_data, order=(5, 1, 0))
model_fit = model.fit()
forecast = model_fit.forecast(steps=len(test_data))
mse = mean_squared_error(test_data, forecast)
rmse = np.sqrt(mse)
print(f'RMSE: {rmse}')
plt.figure(figsize=(12, 6))
 plt.plot(train_data.index, train_data, label='Train Data')
plt.plot(test_data.index, test_data, label='Test Data')
plt.plot(test_data.index, forecast, label='Forecast')
plt.title('Stock Price Prediction')
plt.xlabel('Date')
plt.ylabel('Price (USD)')
plt.legend()
 plt.show()
```





## **RESULT:**

Thus, the program for stock market analysis is executed successfully.

#### **5.LOAN DEFAULT PREDICTION**

**EX.N0:5** 

#### LOAN DEFAULT PREDICTION

**DATE: 21/08/2024** 

**PROBLEM STATEMENT:** Predict loan default probability based on borrower information.

**PYTHON CONCEPTS:** Classes, functions, sequences.

**<u>VISUALIZATION:</u>** ROC curve, bar plots.

**MULTIVARIATE ANALYSIS:** Logistic regression, factor analysis.

**DATASET:** Lending Club Loan Data

#### **ALGORITHM:**

Step 1: Start the program.

Step 2: Import necessary libraries.

Step 3: Load the dataset.

Step 4: Encode categorical variable, define feature & testing set.

Step 5: Split the dataset into training & testing set, create trained model.

Step 6: Print equal metric & test the cell.

#### **PROGRAM:**

import pandas as pd

import matplotlib.pyplot as plt

import seaborn as sns

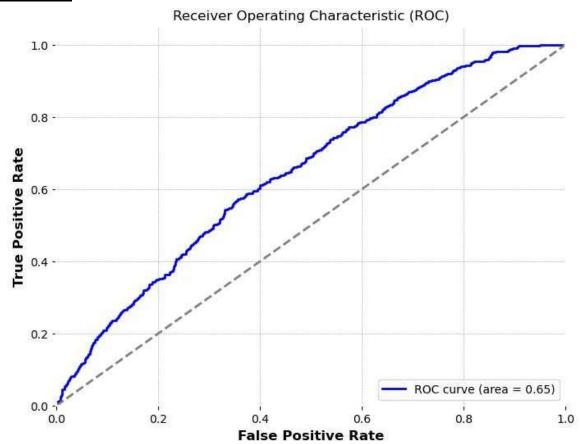
from sklearn.model\_selection import train\_test\_split

from sklearn.linear\_model import LogisticRegression

from sklearn.metrics import roc\_curve, auc

from sklearn.preprocessing import StandardScaler

```
from sklearn.decomposition import PCA
import os
file_path = 'C:/Users/HEMA/Downloads/loan_data.csv' # Update path accordingly
if os.path.exists(file_path):
df = pd.read_csv(file_path)
print("Data loaded successfully.") else:
print(f"File not found: {file_path}")
dummies = pd.get_dummies(df['purpose'], drop_first=True)
df = pd.concat([df, dummies], axis=1)
df.drop('purpose', inplace=True, axis=1)
X = df.drop(['not.fully.paid'], axis=1)
y = df['not.fully.paid']
scaler = StandardScaler()
X_scaled = scaler.fit_transform(X)
pca = PCA(n_components=2)
X_pca = pca.fit_transform(X_scaled)
X_train, X_test, y_train, y_test = train_test_split(X_pca, y, test_size=0.33, random_state=42)
model = LogisticRegression()
model.fit(X_train, y_train)
y_pred_prob = model.predict_proba(X_test)[:, 1]
fpr, tpr, _ = roc_curve(y_test, y_pred_prob)
roc_auc = auc(fpr, tpr)
plt.figure(figsize=(8, 6))
plt.plot(fpr, tpr, color='blue', lw=2, label=f'ROC curve (area = {roc_auc:.2f})')
plt.plot([0, 1], [0, 1], color='gray', linestyle='--')
plt.xlim([0.0, 1.0])
plt.ylim([0.0, 1.05])
plt.xlabel('False Positive Rate')
plt.ylabel('True Positive Rate')
plt.title('Receiver Operating Characteristic (ROC)')
plt.legend(loc='lower right')
plt.show()
```



## **RESULT**:

Thus, the program for loan default prediction is executed successfully.

#### **6. IMAGE CLASSIFICATION**

**EX.N0:6** 

#### **IMAGE CLASSIFICATION**

**DATE: 04/09/2024** 

**PROBLEM STATEMENT:** Classify images into categories using various features.

**PYTHON CONCEPTS:** File handling, classes.

**<u>VISUALIZATION:</u>** Image plots, feature importance plots.

**MULTIVARIATE ANALYSIS:** PCA, clustering.

**DATASET:** CIFAR-10 Dataset

#### **ALGORITHM:**

Step 1: Start the program.

Step 2: Import necessary libraries.

Step 3: Load the dataset.

Step 4: Encode categorical variable, define feature & testing set.

Step 5: Split the dataset into training & testing set, create trained model.

Step 6: Print equal metric & test the cell.

#### **PROGRAM:**

import tensorflow as tf

from tensorflow.keras import layers, models

from tensorflow.keras.preprocessing.image import ImageDataGenerator

import matplotlib.pyplot as plt

import numpy as np

(X\_train, y\_train), (X\_test, y\_test) = tf.keras.datasets.cifar10.load\_data()

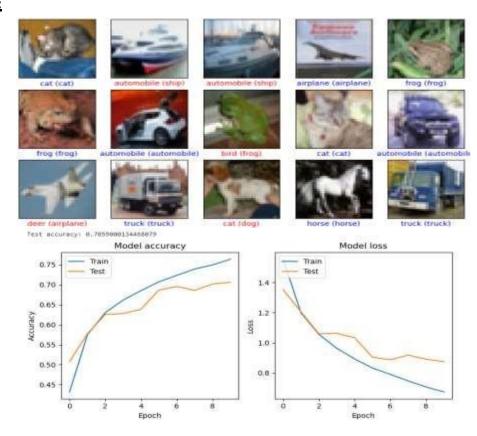
 $X_{train}$ ,  $X_{test} = X_{train} / 255.0$ ,  $X_{test} / 255.0$ 

class\_names = ['airplane', 'automobile', 'bird', 'cat', 'deer',

'dog', 'frog', 'horse', 'ship', 'truck']

```
plt.figure(figsize=(10,10))
 for i in range(25): plt.subplot(5,5,i+1)
plt.xticks([]) plt.yticks([]) plt.grid(False)
 plt.imshow(X_train[i], cmap=plt.cm.binary)
 plt.xlabel(class_names[y_train[i][0]])
 plt.show() model = models.Sequential([
layers.Conv2D(32, (3, 3), activation='relu', input_shape=(32, 32, 3)),
layers.MaxPooling2D((2, 2)),
layers.Conv2D(64, (3, 3), activation='relu'),
layers.MaxPooling2D((2, 2)),
 layers.Conv2D(64, (3, 3), activation='relu'),
layers.Flatten(), layers.Dense(64, activation='relu'),
 layers.Dense(10) ]) model.compile(optimizer='adam',
loss=tf.keras.losses.SparseCategoricalCrossentropy(from_logits=True),
metrics=['accuracy'])
history = model.fit(X_train, y_train, epochs=10,
 validation_data=(X_test, y_test))
 test_loss, test_acc = model.evaluate(X_test, y_test, verbose=2)
 print(f"\nTest accuracy: {test_acc}")
 plt.figure(figsize=(8, 4))
 plt.subplot(1, 2, 1) plt.plot(history.history['accuracy'])
 plt.plot(history.history['val_accuracy'])
plt.title('Model accuracy')
 plt.ylabel('Accuracy') plt.xlabel('Epoch')
 plt.legend(['Train', 'Test'], loc='upper left')
 plt.subplot(1, 2, 2) plt.plot(history.history['loss'])
 plt.plot(history.history['val_loss'])
 plt.title('Model loss') plt.ylabel('Loss')
plt.xlabel('Epoch')
 plt.legend(['Train', 'Test'], loc='upper left')
 plt.tight_layout() plt.show()
 predictions = model.predict(X_test)
 plt.figure(figsize=(10, 10))
```

```
for i in range(25): plt.subplot(5, 5, i+1)
plt.xticks([]) plt.yticks([]) plt.grid(False)
plt.imshow(X_test[i], cmap=plt.cm.binary)
predicted_label = np.argmax(predictions[i])
true_label = y_test[i][0]
color = 'blue' if predicted_label == true_label else 'red'
plt.xlabel(f"{class_names[predicted_label]} ({class_names[true_label]})", color=color)
plt.show()
```



#### **RESULT**:

Thus, the program for Image Classification is executed successfully.

#### **7, PREDICTING DIABETES**

**EX.N0:7** 

#### PREDICTING DIABETES

**DATE: 11/09/2024** 

**PROBLEM STATEMENT:** Predict the onset of diabetes based on medical measurements.

**PYTHON CONCEPTS:** Data structures, numeric types, functions.

**<u>VISUALIZATION:</u>** Scatter plots, heatmaps.

**MULTIVARIATE ANALYSIS:** Logistic regression, LDA.

**DATASET:** Pima Indians Diabetes Database

#### **ALGORITHM:**

Step 1: Start the program.

Step 2: Import necessary libraries.

Step 3: Load the dataset.

Step 4: Encode categorical variable, define feature & testing set.

Step 5: Split the dataset into training & testing set, create trained model.

Step 6: Print equal metric & test the cell.

#### **PROGRAM:**

import pandas as pd

import seaborn as sns

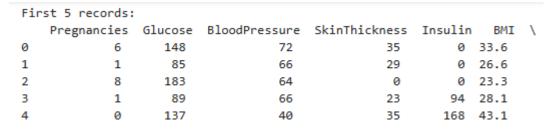
import matplotlib.pyplot as plt

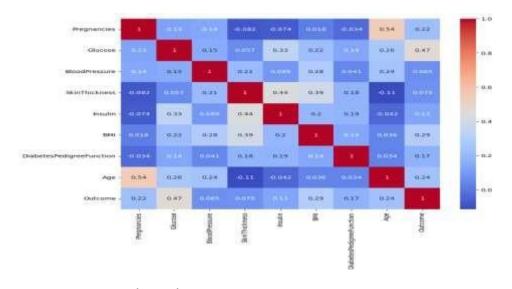
from sklearn.model\_selection import train\_test\_split

from sklearn.linear\_model import LogisticRegression

from sklearn.metrics import classification\_report, confusion\_matrix, accuracy\_score

```
url = https://raw.githubusercontent.com/jbrownlee/Datasets/master/pima-indians-diabetes.data.csv
columns = ['Pregnancies', 'Glucose', 'BloodPressure', 'SkinThickness', 'Insulin', 'BMI',
'DiabetesPedigreeFunction', 'Age', 'Outcome']
data = pd.read_csv(url, header=None, names=columns)
print("First 5 records:\n", data.head())
print("\nStatistical Summary:\n", data.describe())
print("\nDataset Info:\n")
print(data.info())
sns.pairplot(data, hue='Outcome')
plt.show()
correlation_matrix = data.corr()
plt.figure(figsize=(10, 8))
sns.heatmap(correlation_matrix, annot=True, cmap='coolwarm')
plt.show()
X = data.drop('Outcome', axis=1)
y = data['Outcome']
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_state=42)
model = LogisticRegression(max_iter=1000)
model.fit(X_train, y_train)
y_pred = model.predict(X_test)
print("Confusion Matrix:")
print(confusion_matrix(y_test, y_pred))
print("\nClassification Report:")
print(classification_report(y_test, y_pred))
accuracy = accuracy_score(y_test, y_pred)
print(f"\nModel Accuracy: {accuracy * 100:.2f}%")
sample = X_{test.iloc}[0].values.reshape(1, -1)
sample_prediction = model.predict(sample)
print(f"\nPrediction for sample case (1 = Diabetes, 0 = No Diabetes): {sample_prediction[0]}")
```





Confusion Matrix: [[120 31]

[ 30 50]]

Classification Report:

	precision	recall	f1-score	support	
0	0.80	0.79	0.80	151	
1	0.62	0.62	0.62	80	
accuracy			0.74	231	
macro avg weighted avg	0.71 0.74	0.71 0.74	0.71 0.74	231 231	

Model Accuracy: 73.59%

Prediction for sample case (1 = Diabetes, 0 = No Diabetes): 0

## **RESULT**:

Thus, the program for predicting diabetes is executed successfully.

#### **8. WINE OUALITY PREDICTION**

**EX.N0:8** 

## WINE QUALITY PREDICTION

**DATE: 18/09/2024** 

**PROBLEM STATEMENT:** Predict the quality of wine based on various chemical properties.

**PYTHON CONCEPTS:** Classes, sequences, file handling.

**VISUALIZATION:** Histograms, box plots.

**MULTIVARIATE ANALYSIS:** Multiple regression, factor analysis.

**DATASET:** Wine Quality Dataset

#### **ALGORITHM:**

Step 1: Start the program.

Step 2: Import necessary libraries.

Step 3: Load the dataset.

Step 4: Encode categorical variable, define feature & testing set.

Step 5: Split the dataset into training & testing set, create trained model.

Step 6: Print equal metric & test the cell.

#### **PROGRAM:**

import pandas as pd

import numpy as np

import matplotlib.pyplot as plt

import seaborn as sns

from sklearn.model\_selection import train\_test\_split

from sklearn.linear\_model import LinearRegression

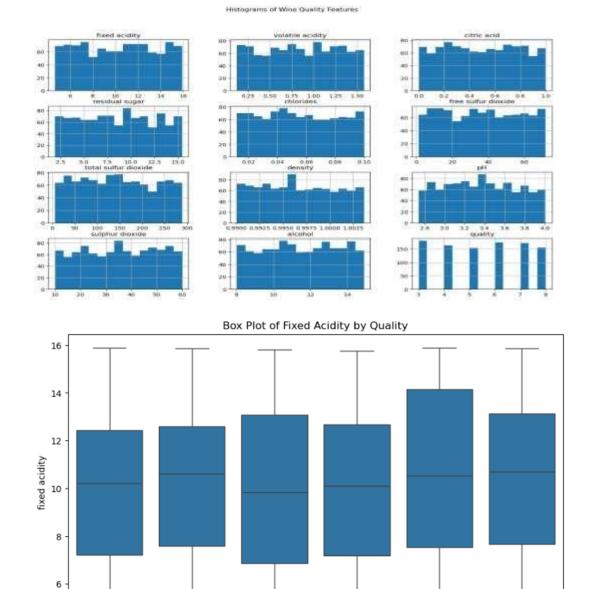
from sklearn.metrics import mean\_squared\_error, r2\_score

class WineQualityPredictor:

def\_\_init\_(self, num\_samples=1000):

```
self.num_samples = num_samples
self.data = None
 self.model = None
 def generate_data(self):
 np.random.seed(42)
 quality = np.random.randint(3, 9, self.num samples) # Quality scores between 3 and 8
fixed_acidity = np.random.uniform(4.6, 15.9, self.num_samples)
 volatile_acidity = np.random.uniform(0.12, 1.58, self.num_samples)
 citric acid = np.random.uniform(0, 1, self.num samples)
 residual sugar = np.random.uniform(1.9, 15.5, self.num samples)
 chlorides = np.random.uniform(0.012, 0.1, self.num_samples)
 free_sulfur_dioxide = np.random.uniform(1, 72, self.num_samples)
 total_sulfur_dioxide = np.random.uniform(6, 289, self.num_samples)
 density = np.random.uniform(0.99007, 1.00369, self.num_samples)
 pH = np.random.uniform(2.74, 4.01, self.num_samples)
 sulfur_dioxide = np.random.uniform(10, 60, self.num_samples)
 alcohol = np.random.uniform(8.0, 14.9, self.num_samples)
 self.data = pd.DataFrame({
 'fixed acidity': fixed_acidity, 'volatile acidity': volatile_acidity, 'citric acid': citric_acid,
 'residual sugar': residual_sugar, 'chlorides': chlorides, 'free sulfur dioxide': free_sulfur_dioxide,
 'total sulfur dioxide': total_sulfur_dioxide, 'density': density, 'pH': pH,
 'sulphur dioxide': sulfur_dioxide, 'alcohol': alcohol, 'quality': quality })
 print(f"Synthetic Data Generated: {self.data.shape[0]} rows and {self.data.shape[1]} columns")
 def visualize_data(self):
 self.data.hist(bins=15, figsize=(15, 10))
 plt.suptitle('Histograms of Wine Quality Features')
 plt.show() plt.figure(figsize=(10, 6))
 sns.boxplot(x='quality', y='fixed acidity', data=self.data)
 plt.title('Box Plot of Fixed Acidity by Quality')
 plt.show() def preprocess_data(self):
 X = self.data.drop('quality', axis=1)
y = self.data['quality']
 return X, y def train_model(self, X, y):
```

```
X train, X test, y train, y test = train test split(X, y, test size=0.2, random state=42)
self.model = LinearRegression()
self.model.fit(X_train, y_train)
y_pred = self.model.predict(X_test)
return y_train, y_test, y_pred
def evaluate_model(self, y_test, y_pred):
mse = mean_squared_error(y_test, y_pred)
r2 = r2\_score(y\_test, y\_pred)
print(f'Mean Squared Error: {mse}') print(f'R^2 Score: {r2}')
def predict_quality(self, input_features):
input df = pd.DataFrame([input features], columns=self.data.columns[:-1])
prediction = self.model.predict(input_df) return prediction[0]
def run(self): self.generate_data() self.visualize_data()
X, y = self.preprocess_data()
y_train, y_test, y_pred = self.train_model(X, y)
self.evaluate_model(y_test, y_pred)
if name == " main ":
wine_predictor = WineQualityPredictor(num_samples=1000)
wine_predictor.run()
example_features = {
'fixed acidity': 7.4, 'volatile acidity': 0.7, 'citric acid': 0.0,
'residual sugar': 1.9, 'chlorides': 0.076, 'free sulfur dioxide': 11.0,
'total sulfur dioxide': 34.0, 'density': 0.9978, 'pH': 3.51,
'sulphur dioxide': 45.0, 'alcohol': 9.4 }
predicted_quality = wine_predictor.predict_quality(example_features)
print(f'Predicted Wine Quality: {predicted_quality:.2f}')
```



5

quality

Mean Squared Error: 2.8525212491984275 R^2 Score: -0.0010251435985495494 Predicted Wine Quality: 5.51

## **RESULT**:

Thus, the program for wine quality prediction is executed successfully.

#### 9. HEART DISEASE PREDICTION

**EX.N0:9** 

#### HEART DISEASE PREDICTION

**DATE: 07/10/2024** 

**PROBLEM STATEMENT:** Predict heart disease based on clinical parameters

**PYTHON CONCEPTS:** Functions, data structures.

**<u>VISUALIZATION:</u>** Pair plots, ROC curve.

**MULTIVARIATE ANALYSIS:** Logistic regression, PCA.

**DATASET:** Heart Disease Dataset

#### **ALGORITHM:**

Step 1: Start the program.

Step 2: Import necessary libraries.

Step 3: Load the dataset.

Step 4: Encode categorical variable, define feature & testing set.

Step 5: Split the dataset into training & testing set, create trained model.

Step 6: Print equal metric & test the cell.

#### **PROGRAM:**

import numpy as np

import pandas as pd

import matplotlib.pyplot as plt

import seaborn as sns

from sklearn.model\_selection import train\_test\_split

from sklearn.preprocessing import StandardScaler

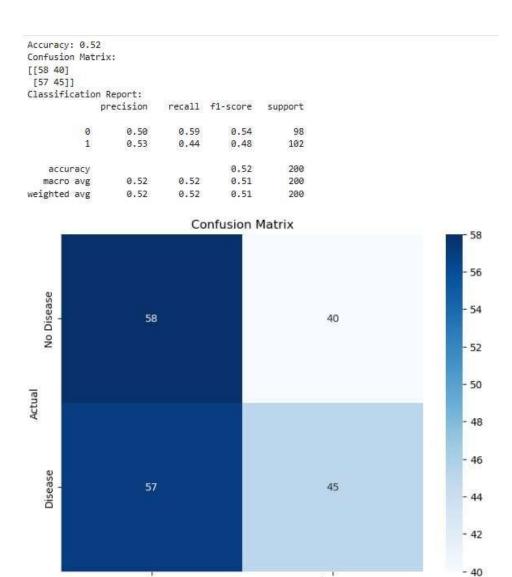
from sklearn.linear\_model import LogisticRegression

 $from \ sklearn.metrics \ import \ accuracy\_score, \ confusion\_matrix, \ classification\_report$ 

np.random.seed(42)

```
num\_samples = 1000
age = np.random.randint(30, 80, num samples)
sex = np.random.randint(0, 2, num_samples)
cp = np.random.randint(0, 4, num_samples)
trestbps = np.random.randint(90, 200, num_samples)
chol = np.random.randint(150, 300, num_samples)
fbs = np.random.randint(0, 2, num_samples)
restecg = np.random.randint(0, 2, num_samples)
thalach = np.random.randint(60, 200, num_samples)
exang = np.random.randint(0, 2, num_samples)
oldpeak = np.random.uniform(0, 6, num_samples)
slope = np.random.randint(0, 3, num_samples)
ca = np.random.randint(0, 4, num_samples)
thal = np.random.randint(1, 4, num_samples)
target = np.random.randint(0, 2, num_samples)
data = pd.DataFrame({
 'age': age, 'sex': sex, 'cp': cp,
'trestbps': trestbps, 'chol': chol,
'fbs': fbs, 'restecg': restecg, 'thalach': thalach, 'exang': exang,
'oldpeak': oldpeak, 'slope': slope, 'ca': ca,
'thal': thal, 'target': target})
X = data.drop('target', axis=1)
y = data['target']
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
scaler = StandardScaler()
X_{train} = scaler.fit_{transform}(X_{train})
X_{\text{test}} = \text{scaler.transform}(X_{\text{test}})
model = LogisticRegression()
model.fit(X_train, y_train)
y_pred = model.predict(X_test)
accuracy = accuracy score(y test, y pred)
conf_matrix = confusion_matrix(y_test, y_pred)
class_report = classification_report(y_test, y_pred)
```

```
print(f'Accuracy: {accuracy:.2f}')
print('Confusion Matrix:')
print(conf_matrix)
print('Classification Report:')
print(class_report)
plt.figure(figsize=(8, 6))
sns.heatmap(conf_matrix, annot=True, fmt='d', cmap='Blues', xticklabels=['No Disease',
'Disease'], yticklabels=['No Disease', 'Disease'])
plt.title('Confusion Matrix')
plt.xlabel('Predicted')
plt.ylabel('Actual')
plt.show()
importance = model.coef_[0]
features = X.columns
importance_df = pd.DataFrame({'Feature': features, 'Importance': importance})
importance_df = importance_df.sort_values(by='Importance', ascending=False)
plt.figure(figsize=(10, 6))
sns.barplot(data=importance_df, x='Importance', y='Feature', palette='viridis')
plt.title('Feature Importance')
plt.xlabel('Coefficient Value')
plt.ylabel('Features')
plt.axvline(0, color='red', linestyle='--') # Adding a vertical line at 0
plt.show()
```



Predicted

Disease

## **RESULT**:

Thus, the program for heart disease prediction is executed successfully.

No Disease

#### 10. BREAST CANCER DIAGNOSIS

EX.N0: 10	Procest Consen Diagnosis
DATE: 09/10/2024	Breast Cancer Diagnosis

**PROBLEM STATEMENT:** Classify tumors as benign or malignant based on features.

**PYTHON CONCEPTS:** Classes, sequences.

**<u>VISUALIZATION:</u>** Confusion matrix, bar plots.

**MULTIVARIATE ANALYSIS:** LDA, logistic regression.

**DATASET:** Breast Cancer Wisconsin Dataset

#### **ALGORITHM:**

Step 1: Start the program.

Step 2: Import necessary libraries.

Step 3: Load the dataset.

Step 4: Encode categorical variable, define feature & testing set.

Step 5: Split the dataset into training & testing set, create trained model.

Step 6: Print equal metric & test the cell.

#### **PROGRAM:**

import numpy as np

import pandas as pd

import matplotlib.pyplot as plt

import seaborn as sns

from sklearn.model\_selection import train\_test\_split

from sklearn.preprocessing import StandardScaler

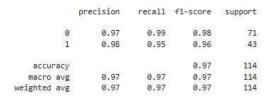
from sklearn.linear\_model import LogisticRegression

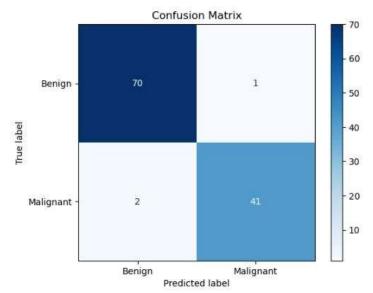
 $from \ sklearn.metrics \ import \ accuracy\_score, \ confusion\_matrix, \ classification\_report$ 

np.random.seed(42)

```
num\_samples = 1000
age = np.random.randint(30, 80, num samples)
sex = np.random.randint(0, 2, num_samples)
cp = np.random.randint(0, 4, num_samples)
trestbps = np.random.randint(90, 200, num_samples)
chol = np.random.randint(150, 300, num_samples)
fbs = np.random.randint(0, 2, num_samples)
restecg = np.random.randint(0, 2, num_samples)
thalach = np.random.randint(60, 200, num_samples)
exang = np.random.randint(0, 2, num_samples)
oldpeak = np.random.uniform(0, 6, num_samples)
slope = np.random.randint(0, 3, num_samples)
ca = np.random.randint(0, 4, num_samples)
thal = np.random.randint(1, 4, num_samples)
target = np.random.randint(0, 2, num_samples)
data = pd.DataFrame({
 'age': age, 'sex': sex, 'cp': cp,
'trestbps': trestbps, 'chol': chol,
'fbs': fbs, 'restecg': restecg, 'thalach': thalach, 'exang': exang,
'oldpeak': oldpeak, 'slope': slope, 'ca': ca,
'thal': thal, 'target': target})
X = data.drop('target', axis=1)
y = data['target']
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
scaler = StandardScaler()
X_{train} = scaler.fit_{transform}(X_{train})
X_{\text{test}} = \text{scaler.transform}(X_{\text{test}})
model = LogisticRegression()
model.fit(X_train, y_train)
y_pred = model.predict(X_test)
accuracy = accuracy score(y test, y pred)
conf_matrix = confusion_matrix(y_test, y_pred)
class_report = classification_report(y_test, y_pred)
```

```
print(f'Accuracy: {accuracy:.2f}')
print('Confusion Matrix:')
print(conf_matrix)
print('Classification Report:')
print(class_report)
plt.figure(figsize=(8, 6))
sns.heatmap(conf_matrix, annot=True, fmt='d', cmap='Blues', xticklabels=['No Disease',
'Disease'], yticklabels=['No Disease', 'Disease'])
plt.title('Confusion Matrix')
plt.xlabel('Predicted')
plt.ylabel('Actual')
plt.show()
importance = model.coef_[0]
features = X.columns
importance_df = pd.DataFrame({'Feature': features, 'Importance': importance})
importance_df = importance_df.sort_values(by='Importance', ascending=False)
plt.figure(figsize=(10, 6))
sns.barplot(data=importance_df, x='Importance', y='Feature', palette='viridis')
plt.title('Feature Importance')
plt.xlabel('Coefficient Value')
plt.ylabel('Features')
plt.axvline(0, color='red', linestyle='--') # Adding a vertical line at 0
plt.show()
```





Enter the following features for prediction: compactness\_se: 0.03 concavity\_se: 0.03 radius\_mean: 14.5 concave points\_se: 0.02 texture\_mean: 20.0 symmetry\_se: 0.02 perimeter\_mean: 90.0 fractal\_dimension\_se: 0.003 area\_mean: 560.0 radius\_worst: 16.0 smoothness\_mean: 0.1 texture\_worst: 25.0 compactness\_mean: 0.15 perimeter\_worst: 100.0 concavity\_mean: 0.2 area\_worst: 800.0 concave points\_mean: 0.1 smoothness\_worst: 0.14 symmetry\_mean: 0.18 compactness\_worst: 0.25 fractal\_dimension\_mean: 0.06 concavity\_worst: 0.3 radius\_se: 0.6 concave points\_worst: 0.15 texture\_se: 1.2 symmetry\_worst: 0.25 perimeter\_se: 10.0 fractal\_dimension\_worst: 0.08 The tumor is predicted to be: Malignant area\_se: 40.0 Based on the symptoms provided, the person may be at risk. smoothness\_se: 0.007

#### **RESULT:**

Thus, the program for breast cancer diagnosis is executed successfully.

#### 11. PREDICTING FLIGHT DELAYS

EX.N0:11

#### PREDICTING FLIGHT DELAYS

**DATE:** 16/10/2024

**PROBLEM STATEMENT:** Predict flight delays based on historical data.

**PYTHON CONCEPTS:** File reading/writing, functions.

**<u>VISUALIZATION:</u>** Line plots, scatter plots.

**MULTIVARIATE ANALYSIS:** Regression, clustering.

**DATASET:** Flight Delay Dataset

#### **ALGORITHM:**

Step 1: Start the program.

Step 2: Import necessary libraries.

Step 3: Load the dataset.

Step 4: Encode categorical variable, define feature & testing set.

Step 5: Split the dataset into training & testing set, create trained model.

Step 6: Print equal metric & test the cell.

#### **PROGRAM:**

import pandas as pd

import numpy as np

import matplotlib.pyplot as plt

import seaborn as sns

from sklearn.model\_selection import train\_test\_split

from sklearn.linear\_model import LinearRegression

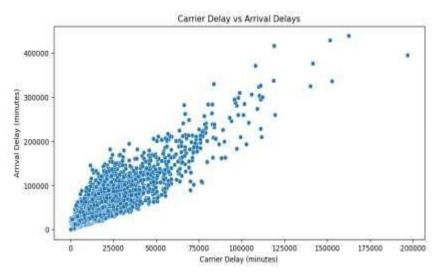
from sklearn.metrics import mean\_absolute\_error, mean\_squared\_error, r2\_score

df = pd.read\_csv('C:/Users/HEMA/Downloads/Airline\_Delay\_Cause.csv')

print(df.columns)

```
print(df.isnull().sum())
 df.dropna(inplace=True) # or df.fillna(method='ffill', inplace=True)
 if 'year' in df.columns and 'month' in df.columns:
 df['date'] = pd.to_datetime(df[['year', 'month']].assign(day=1))
 plt.figure(figsize=(10, 5))
 sns.lineplot(data=df, x='date', y='arr_delay') # Adjust if necessary
 plt.title('Flight Delays Over Time')
 plt.xticks(rotation=45)
 plt.show()
 delay column = 'arr delay' # Using 'arr delay' for now
 if 'carrier_delay' in df.columns and delay_column in df.columns:
 plt.figure(figsize=(10, 5))
 sns.scatterplot(data=df, x='carrier_delay', y=delay_column) # Adjust as needed
 plt.title('Carrier Delay vs Arrival Delays') plt.xlabel('Carrier Delay (minutes)')
 plt.ylabel('Arrival Delay (minutes)') plt.show()
 else: print("Check the delay columns: 'carrier_delay' or 'arr_delay' do not exist in the
 DataFrame.")
 df['day_of_week'] = df['date'].dt.dayofweek # Monday=0, Sunday=6
 features = ['day_of_week', 'arr_flights', 'carrier_ct'] # Modify as needed
 X = df[features] y = df[delay\_column]
 X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
 model = LinearRegression()
 model.fit(X_train, y_train)
 predictions = model.predict(X_test)
 print('Mean Absolute Error:', mean_absolute_error(y_test, predictions))
 print('Mean Squared Error:', mean_squared_error(y_test, predictions))
 print('R-squared:', r2_score(y_test, predictions))
 plt.figure(figsize=(10, 5)) plt.scatter(y_test, predictions)
 plt.plot([min(y_test), max(y_test)], [min(y_test), max(y_test)], color='red', linewidth=2) # Line
 of equality
 plt.title('Predictions vs Actual Delays') plt.xlabel('Actual Delays')
 plt.ylabel('Predicted Delays') plt.show()
```

```
'security_delay', 'late_aircraft_delay'],
     dtype='object')
year
month
                      0
carrier
                      0
carrier_name
                      0
                                                   Flight Delays Over Time
                          10000
airport
                      0
                      0
airport_name
arr_flights
                    240
                           8000
arr_del15
                    443
carrier_ct
                    240
weather_ct
                    240
nas_ct
                    240
security_ct
                    240
                    240
late_aircraft_ct
arr_cancelled
                    240
arr_diverted
                    240
                           2000
arr_delay
                    240
carrier_delay
                    240
weather_delay
                    240
nas delay
                    240
security_delay
                    240
late_aircraft_delay
                    240
dtype: int64
```



Mean Absolute Error: 1592.2201262853362 Mean Squared Error: 25524907.35571326

R-squared: 0.8439698040165798

#### **RESULT**:

Thus, the program for predicting flight delays is executed successfully.

#### 12. ENERGY CONSUMPTION FORECASTING

EX.N0:12

## **ENERGY CONSUMPTION FORECASTING**

**DATE: 23/10/2024** 

**PROBLEM STATEMENT:** Forecast energy consumption based on historical data.

**PYTHON CONCEPTS:** Functions, numeric types.

**VISUALIZATION:** Line plots, heatmaps.

**MULTIVARIATE ANALYSIS:** Time series analysis, regression.

**DATASET:** Energy Consumption Dataset

#### **ALGORITHM:**

Step 1: Start the program.

Step 2: Import necessary libraries.

Step 3: Load the dataset.

Step 4: Encode categorical variable, define feature & testing set.

Step 5: Split the dataset into training & testing set, create trained model.

Step 6: Print equal metric & test the cell.

#### **PROGRAM:**

import pandas as pd

import numpy as np

import matplotlib.pyplot as plt

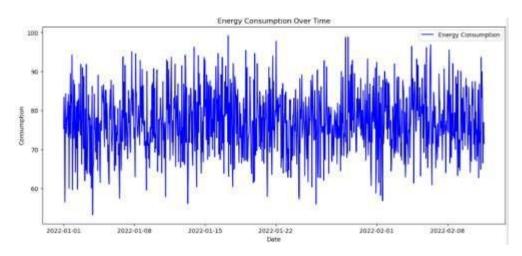
import seaborn as sns

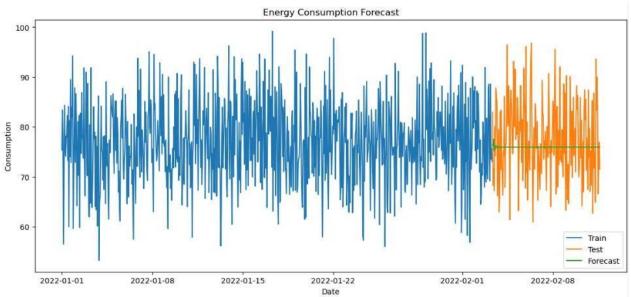
from statsmodels.tsa.arima.model import ARIMA

from sklearn.metrics import mean\_squared\_error

```
data =
pd.read_csv('C:/Users/HEMA/Downloads/energy_consumption_dataset.csv',
parse_dates=['Timestamp'], index_col='Timestamp')
print(data.head()) print(data.info())
data = data.fillna(method='ffill')
plt.figure(figsize=(14, 6))
plt.plot(data['EnergyConsumption'], color='blue', label='Energy Consumption')
plt.title('Energy Consumption Over Time')
plt.xlabel('Date') plt.ylabel('Consumption')
plt.legend() plt.show()
numeric_data = data.select_dtypes(include=[np.number])
plt.figure(figsize=(10, 8))
sns.heatmap(numeric_data.corr(), annot=True, cmap='coolwarm')
plt.title('Correlation Matrix') plt.show()
from statsmodels.tsa.seasonal import seasonal decompose
result = seasonal_decompose(data['EnergyConsumption'], model='additive', period=24) # Adjust
period based on your data's frequency
result.plot() plt.show()
train\_size = int(len(data) * 0.8)
train, test = data['EnergyConsumption'][:train size], data['EnergyConsumption'][train size:]
model = ARIMA(train, order=(5, 1, 0)) # Adjust (p,d,q) based on your data's behavior
fitted_model = model.fit()
forecast = fitted model.forecast(steps=len(test))
forecast index = test.index
mse = mean_squared_error(test, forecast)
rmse = np.sqrt(mse)
print(f'RMSE: {rmse}')
plt.figure(figsize=(14, 6))
plt.plot(train, label='Train')
plt.plot(test, label='Test')
plt.plot(forecast_index, forecast, label='Forecast')
plt.title('Energy Consumption Forecast')
plt.xlabel('Date')
plt.ylabel('Consumption')
plt.legend()
plt.show()
```

	Temperature	Humidity	SquareFootage	Occupancy	V		HVACUsage	LightingUsage	RenewableEnergy	DayOfWeek
Timestamp						Timestamp				
2022-01-01 00:00:00	25.139433	43.431581	1565.693999	5		2022-01-01 00:00:00	0 On	Off	2.774699	Monday
2022-01-01 01:00:00	27.731651	54.225919	1411.064918	1		2022-01-01 01:00:00	0n	On-	21.831384	Saturday
2022-01-01 02:00:00	28.704277	58.907658	1755.715009	2		2022-01-01 02:00:00	0ff	Off	6.764672	Sunday
2022-01-01 03:00:00	20.080469	50.371637	1452.316318	1		2022-01-01 03:00:00	0ff	On	8.623447	Wednesday
2022-01-01 04:00:00	23.097359	51.401421	1094,130359	9		2022-01-01 04:00:00	) On	Off	3.071969	Friday





## **RESULT**:

Thus, the program for energy consumption forecasting is executed successfully.