| EX.N0:1 | |
|---------|-------------------------|
| DATE: | PREDICTING HOUSE PRICES |

<u>AIM:</u> To build a regression model to predict house prices based on features like location, size, and amenities.

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

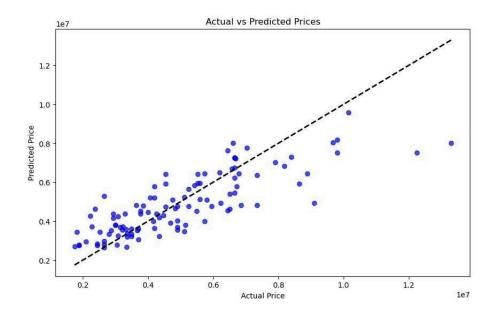
import r2_score, mean_absolute_error

import matplotlib.pyplot as plt

file path = 'C:/Users/APPU/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') plt.show() print(f'R-squared (R<sup>2</sup>): {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.

| EX.N0:2 | CUSTOMER SEGMENTATION FOR AN |
|---------|------------------------------|
| DATE: | E-COMMERCE COMPANY |

To perform cluster analysis to segment customers based on purchasing behaviour.

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

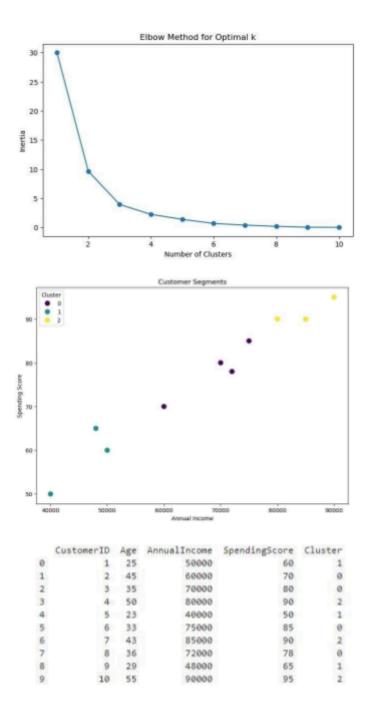
import seaborn as sns

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.

| EX.N0:3 | SENTIMENT ANALYSIS OF MOVIE REVIEWS |
|---------|-------------------------------------|
| DATE: | |

To classify movie reviews as positive or negative using text 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

sklearn.model selection import train test split

from sklearn.preprocessing import LabelEncoder

import nltk from nltk.corpus

import stopwords

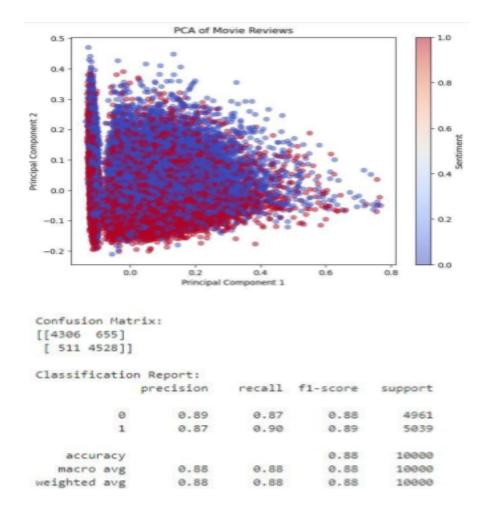
from nltk.tokenize import word tokenize

from nltk.stem import PorterStemmer

import seaborn as sns nltk.download('punkt') nltk.download('stopwords')

```
df = pd.read csv('C:/Users/AI LAB/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()
```



RESULT:

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

| EX.NO : 4 | HANDWRITTEN DIGIT RECOGNITION |
|-----------|-------------------------------|
| DATE: | |

To implement handwritten digit recognition using python.

ALGORITHM:

Step 1: Start the program.

Step 2: Import necessary libraries.

Step 3: Load the dataset.

Step 4: Extract and define feature in testing set and training set.

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

Step 6: Print equal metric & test the cell.

SOURCE CODE:

import numpy as np

from keras.datasets import mnist

from keras.utils import to categorical

from keras.models import Sequential

from keras.layers import Input, Flatten, Dense

import matplotlib.pyplot as plt

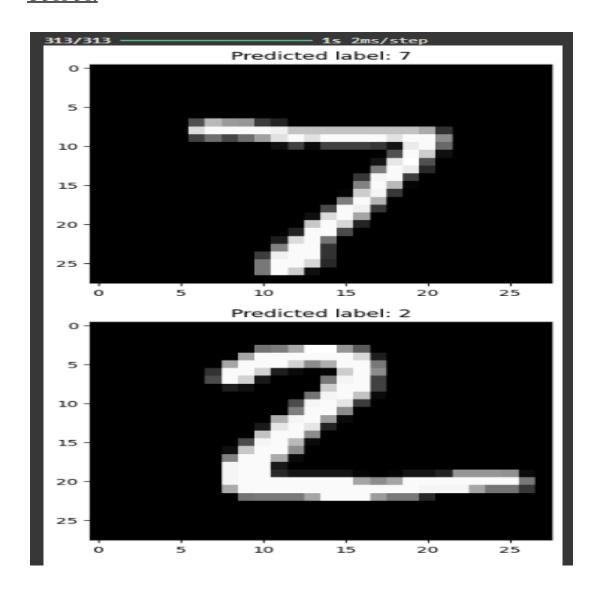
(x train, y train), (x test, y test) = mnist.load data()

plt.imshow(x train[0], cmap='gray')

plt.show()

```
x train = x train.astype('float32') / 255.0
x test = x test.astype('float32') / 255.0
y train = to categorical(y train, num classes=10)
y test = to categorical(y test, num classes=10)
# Define the model
model = Sequential([
  Input(shape=(28, 28)),
  Flatten(),
  Dense(128, activation='relu'),
  Dense(128, activation='relu'),
  Dense(10, activation='softmax')
])
model.compile(optimizer='adam', loss='categorical crossentropy', metrics=['accuracy'])
model.summary()
model.fit(x_train, y_train, epochs=10)
test loss, test acc = model.evaluate(x test, y test)
print(f'Test accuracy: {test acc}')
predictions = model.predict(x test)
for i in range(5):
  plt.imshow(x test[i], cmap='gray')
```

 $plt.title(f'Predicted\ label:\ \{np.argmax(predictions[i])\}')$ plt.show()



RESULT:

Thus the program for implementing handwritten digit recognition is successfully executed and output is obtained.

| EX.N0:5 | STOCK MARKET ANALYSIS |
|---------|-----------------------|
| DATE: | |

To analyse stock market data to predict future stock 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

import matplotlib.pyplot as plt

import mplfinance as mpf

from statsmodels.tsa.arima.model

import ARIMA

from sklearn.metrics import mean squared error

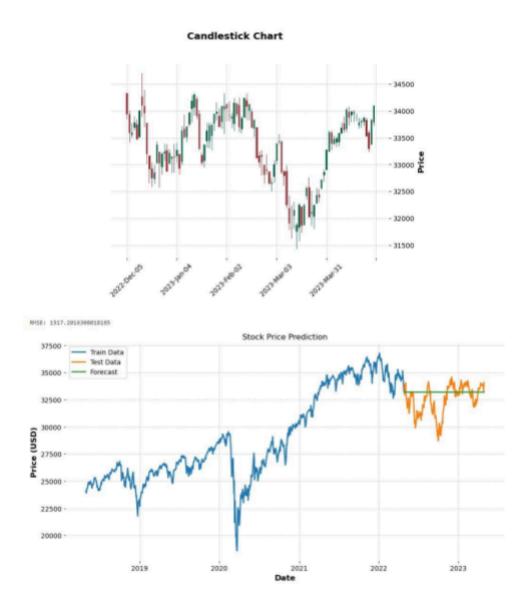
import numpy as np

 $file path = r'C: \label{eq:conditional} When the path = r'C: \la$

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:]
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.

| EX.N0:6 | LOAN DEFAULT PREDICTION |
|---------|-------------------------|
| DATE: | |

Predict loan default probability based on borrower information.

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

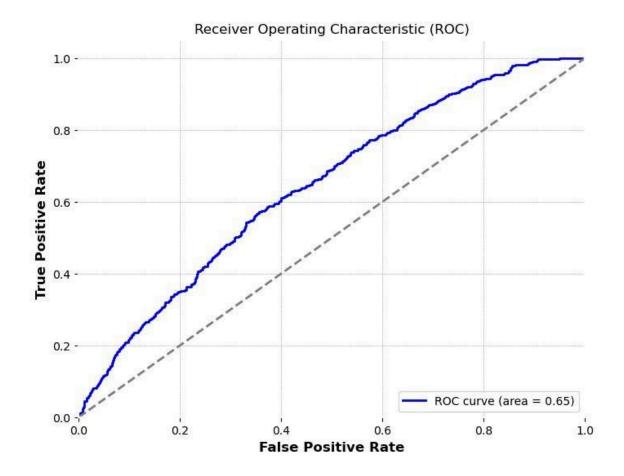
import StandardScaler from sklearn.decomposition

import PCA import os

file path = 'C:/Users/APPU/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.

| EX.NO: 7 | IMAGE RECOGNITION |
|----------|-------------------|
| DATE: | IWAGE RECOGNITION |

To implement image recognition to classify images into categories using various features.

ALGORITHM:

Step 1: Start the program.

Step 2: Import necessary libraries.

Step 3: Load the dataset.

Step 4: Extract and define feature in testing set and training set.

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

Step 6: Print equal metric & test the cell.

SOURCE CODE:

import numpy as np

import matplotlib.pyplot as plt

from sklearn.decomposition import PCA

from sklearn.cluster import KMeans

from keras.datasets import cifar10

from sklearn.preprocessing import StandardScaler

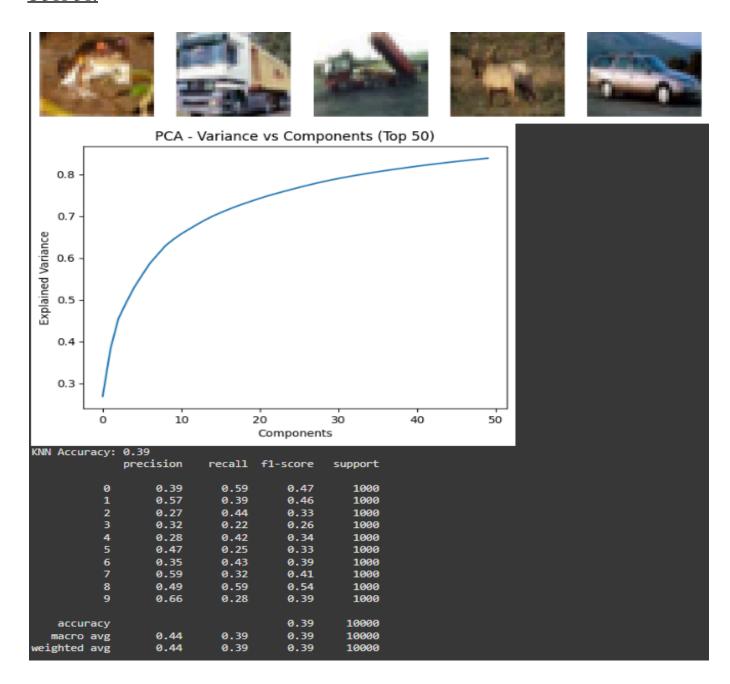
from sklearn.neighbors import KNeighborsClassifier

from sklearn.metrics import accuracy_score, classification report

```
class SimpleImageClassifier:
  def __init (self):
    (self.X train, self.y train), (self.X test, self.y test) = cifar10.load data()
    self.X train = self.X train.reshape(len(self.X train), -1) / 255.0
    self.X test = self.X test.reshape(len(self.X test), -1) / 255.0
  def visualize images(self, n=5):
    plt.figure(figsize=(10, 2))
     for i in range(n):
       plt.subplot(1, n, i + 1)
       plt.imshow(self.X train[i].reshape(32, 32, 3))
       plt.axis('off')
    plt.show()
  def pca analysis(self, components=50):
    scaler = StandardScaler()
    self.X train scaled = scaler.fit transform(self.X train)
    self.X test scaled = scaler.transform(self.X test)
    pca = PCA(n components=components)
    self.X train pca = pca.fit transform(self.X train scaled)
    self.X test pca = pca.transform(self.X test scaled)
    cumulative explained variance = np.cumsum(pca.explained variance ratio )
    plt.plot(cumulative explained variance)
```

```
plt.xlabel('Number of Components')
    plt.ylabel('Cumulative Explained Variance')
    plt.title(fPCA - Variance vs Components (Top {components})')
    plt.show()
  def train knn(self, n neighbors=5):
    knn = KNeighborsClassifier(n neighbors=n neighbors)
    knn.fit(self.X train pca, self.y train.ravel())
    y pred = knn.predict(self.X test pca)
    accuracy = accuracy score(self.y test, y pred)
    print(f"KNN Accuracy with {n neighbors} neighbors: {accuracy:.2f}")
    print(classification report(self.y test, y pred))
  def clustering(self, n clusters=10):
    kmeans = KMeans(n clusters=n clusters, random state=42)
    cluster labels = kmeans.fit predict(self.X train pca)
    plt.scatter(self.X_train_pca[:, 0], self.X_train_pca[:, 1], c=cluster_labels, cmap='rainbow', s=2)
    plt.title(f'KMeans Clustering on CIFAR-10 Images (PCA Reduced, {n clusters})')
    plt.show()
if name == " main ":
  classifier = SimpleImageClassifier()
  classifier.visualize images()
```

classifier.pca_analysis(components=50)
classifier.train_knn(n_neighbors=5)
classifier.clustering(n_clusters=10)



RESULT:

Thus the program for implementing image recognition to classify images into categories using various features is successfully executed and output is obtained.

| EX.NO: 8 | PREDICTING DIABETES |
|----------|---------------------|
| DATE: | |

To predict the onset of diabetes based on medical measurements.

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.

SOURCE CODE:

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.preprocessing import StandardScaler

from sklearn.linear model import LogisticRegression

from sklearn.discriminant analysis import LinearDiscriminantAnalysis

```
from sklearn.metrics import confusion matrix, classification report
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(data.head())
print(data.describe())
sns.pairplot(data, hue='Outcome')
plt.show()
plt.figure(figsize=(10, 8))
sns.heatmap(data.corr(), annot=True, fmt=".2f", cmap="coolwarm")
plt.title("Correlation Heatmap")
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=0.2, random state=42)
scaler = StandardScaler()
X train scaled = scaler.fit transform(X train)
X test scaled = scaler.transform(X test)
log reg = LogisticRegression()
```

```
log_reg.fit(X_train_scaled, y_train)

y_pred_log = log_reg.predict(X_test_scaled)

print("Logistic Regression Classification Report:")

print(classification_report(y_test, y_pred_log))

print("Confusion Matrix:\n", confusion_matrix(y_test, y_pred_log))

lda = LinearDiscriminantAnalysis()

lda.fit(X_train_scaled, y_train)

y_pred_lda = lda.predict(X_test_scaled)

print("LDA Classification Report:")

print(classification_report(y_test, y_pred_lda))

print("Confusion Matrix:\n", confusion_matrix(y_test, y_pred_lda))
```

| LDA Classific | ation Report: | | | |
|---------------------------------------|---------------|--------|----------|---------|
| | precision | recall | f1-score | support |
| 0 | 0.82 | 0.81 | 0.81 | 99 |
| 1 | 0.66 | 0.67 | 0.67 | 55 |
| accuracy | | | 0.76 | 154 |
| macro avg | 0.74 | 0.74 | 0.74 | 154 |
| weighted avg | 0.76 | 0.76 | 0.76 | 154 |
| Confusion Mat [[80 19] [18 37]] | rix: | | | |

RESULT:

Thus the program to predict the onset of diabetes based on medical measurements is successfully predicted and output is obtained.

| EX.NO : 9 | WINE QUALITY PREDICTION |
|-----------|-------------------------|
| DATE: | |

To predict the quality of wine based on various chemical properties.

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.

SOURCE CODE:

import pandas as pd

import numpy as np

import seaborn as sns

import matplotlib.pyplot as plt

from sklearn.model_selection import train_test_split

from sklearn.linear model import LinearRegression

from sklearn.metrics import mean squared error, r2 score

from sklearn.decomposition import FactorAnalysis

```
import pickle
url = "https://archive.ics.uci.edu/ml/machine-learning-databases/wine-quality/winequality-red.csv"
wine data = pd.read csv(url, sep=";")
print(wine data.head())
print(wine data.isnull().sum())
plt.figure(figsize=(8,6))
sns.histplot(wine data['quality'], bins=10, kde=True)
plt.title('Distribution of Wine Quality')
plt.xlabel('Wine Quality')
plt.ylabel('Frequency')
plt.show()
plt.figure(figsize=(16,10))
sns.boxplot(data=wine_data, orient='h')
plt.title('Boxplot of Wine Chemical Properties')
plt.show()
X = wine data.drop('quality', axis=1)
y = wine data['quality']
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_state=42)
regressor = LinearRegression()
regressor.fit(X train, y train)
```

```
y_pred = regressor.predict(X_test)
mse = mean squared error(y test, y pred)
r2 = r2 score(y test, y pred)
print(f"Mean Squared Error: {mse}")
print(f"R-Squared: {r2}")
fa = FactorAnalysis(n_components=6, random state=42)
X fa = fa.fit transform(X)
X train fa, X test fa, y train fa, y test fa = train test split(X fa, y, test size=0.3, random state=42)
regressor fa = LinearRegression()
regressor fa.fit(X train fa, y train fa)
y pred fa = regressor fa.predict(X test fa)
mse fa = mean squared error(y test fa, y pred fa)
r2 fa = r2 score(y test fa, y pred fa)
print(f"Factor Analysis - Mean Squared Error: {mse fa}")
print(f"Factor Analysis - R-Squared: {r2 fa}")
class WineQualityModel:
  def init (self, model):
     self.model = model
  def save model(self, filename='wine quality model.pkl'):
     with open(filename, 'wb') as file:
       pickle.dump(self.model, file)
```

```
print(f"Model saved to {filename}")

def load_model(self, filename='wine_quality_model.pkl'):
    with open(filename, 'rb') as file:
        self.model = pickle.load(file)

    print(f"Model loaded from {filename}")

def predict_quality(self, X):
    return self.model.predict(X)

wine_quality_model = WineQualityModel(regressor)

wine_quality_model.save_model()

wine_quality_model.load_model()

predictions = wine_quality_model.predict_quality(X_test)

print(predictions[:5])
```

Model saved to wine_quality_model.pkl
Model loaded from wine_quality_model.pkl
[5.35676319 5.09071476 5.62553757 5.44886088 5.74478368]

RESULT:

Thus the program to predict the quality of wine based on various chemical properties is successfully executed and output is obtained.

| EX.NO: 10 | HEART DISEASE PREDICTION |
|-----------|--------------------------|
| DATE: | |

To predict heart disease based on clinical parameters.

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.

SOURCE CODE:

import pandas as pd

import numpy as np

import seaborn as sns

import matplotlib.pyplot as plt

from sklearn.model selection import train test split

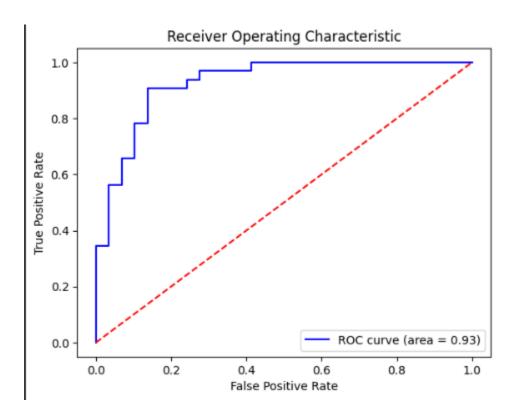
from sklearn.linear model import LogisticRegression

from sklearn.metrics import confusion matrix, roc curve, auc

```
from sklearn.decomposition import PCA
data = pd.read csv("/content/dataset.csv")
print(data.info())
sns.pairplot(data, hue='target')
plt.show()
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)
model = LogisticRegression()
model.fit(X train, y train)
y pred = model.predict(X test)
y probs = model.predict proba(X test)[:, 1]
conf matrix = confusion matrix(y test, y pred)
fpr, tpr, thresholds = roc curve(y test, y probs)
roc auc = auc(fpr, tpr)
print("Confusion Matrix:")
print(conf matrix)
plt.plot(fpr, tpr, color='blue', label='ROC curve (area = %0.2f)' % roc auc)
plt.plot([0, 1], [0, 1], color='red', linestyle='--')
plt.xlabel('False Positive Rate')
plt.ylabel('True Positive Rate')
```

plt.title('Receiver Operating Characteristic')

plt.legend(loc='lower right') plt.show()



RESULT:

Thus the program to predict heart disease based on clinical parameters is successfully executed and output is obtained.

| EX.N0:11 | BREAST CANCER DIAGNOSIS |
|----------|-------------------------|
| DATE: | |

To classify tumors as benign or malignant based on features.

ALGORITHM:

Step 1: Start the program.

Step 2: Import necessary libraries.

Step 3: Load the dataset.

Step 4: Extract and define feature in testing set and training set.

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

Step 6: Print equal metric & test the cell.

SOURCE CODE:

import pandas as pd

import numpy as np

import seaborn as sns

import matplotlib.pyplot as plt

from sklearn.model_selection import train_test_split

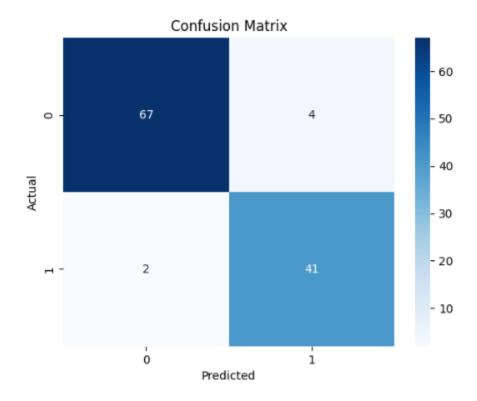
from sklearn.linear_model import LogisticRegression

from sklearn.metrics import confusion_matrix, accuracy_score, classification_report

```
from sklearn.discriminant analysis import LinearDiscriminantAnalysis
data = pd.read csv('/content/breast cancer.csv')
print(data.info())
print(data.describe())
X = data.drop(['id', 'diagnosis'], axis=1)
y = data['diagnosis'].map(\{'M': 1, 'B': 0\})
X train, X test, y train, y test = train test split(X, y, test size=0.2, random state=42)
lda = LinearDiscriminantAnalysis()
X train lda = lda.fit transform(X train, y train)
X test Ida = Ida.transform(X test)
model = LogisticRegression(max iter=1000)
model.fit(X_train_lda, y_train)
y pred = model.predict(X test lda)
conf matrix = confusion matrix(y_test, y_pred)
accuracy = accuracy score(y test, y pred)
class report = classification report(y test, y pred)
sns.heatmap(conf matrix, annot=True, fmt='d', cmap='Blues')
plt.ylabel('Actual')
plt.xlabel('Predicted')
plt.title('Confusion Matrix')
plt.show()
```

print(f'Accuracy: {accuracy:.2f}')

print(class_report)



| Accuracy: 0.95 precision recall f1-score support | | | | | |
|---|--------------|--------------|----------------------|-------------------|--|
| 0 1 | 0.97 0.91 | 0.94 0.95 | 0.96 0.93 | 71 43 | |
| accuracy macro avg weighted avg | 0.94 0.95 | 0.95 0.95 | 0.95 0.94 0.95 | 114 114 114 | |

RESULT:

Thus the classification of tumors as benign or malignant is successfully executed and output is obtained.

| EX.N0:12 | PREDICTING FLIGHT DELAYS |
|----------|--------------------------|
| DATE: | |

To predict flight delays based on historical 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.

SOURCE CODE:

Import necessary libraries

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.cluster import KMeans
from sklearn.metrics import mean squared error, r2 score
data = pd.read csv('flight delay dataset.csv') # Replace with your file path
print("Dataset Head:")
print(data.head())
print("\nDataset Information:")
print(data.info())
print("\nDataset Statistics:")
print(data.describe())
plt.figure(figsize=(10, 6))
plt.plot(data['Date'], data['Delay'], label='Delay over Time')
plt.xlabel('Date')
plt.ylabel('Delay (Minutes)')
plt.title('Flight Delay Trend Over Time')
plt.xticks(rotation=45)
plt.legend()
plt.show()
plt.figure(figsize=(8, 6))
sns.scatterplot(x='Distance', y='Delay', data=data, hue='Airline', palette='viridis')
plt.xlabel('Distance (Miles)')
plt.ylabel('Delay (Minutes)')
```

```
plt.title('Flight Delay vs Distance')
plt.show()
X = data[['Distance', 'Airline_Code', 'Scheduled_Departure_Hour']]
y = data['Delay']
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
regressor = LinearRegression()
regressor.fit(X_train, y_train)
y_pred = regressor.predict(X_test)
mse = mean_squared_error(y_test, y_pred)
r2 = r2\_score(y\_test, y\_pred)
print(f"\nMean Squared Error: {mse}")
print(f"R-squared: {r2}")
plt.figure(figsize=(8, 6))
plt.scatter(y_test, y_pred, color='blue', alpha=0.5)
plt.plot([y.min(), y.max()], [y.min(), y.max()], 'r--')
plt.xlabel('Actual Delay')
plt.ylabel('Predicted Delay')
plt.title('Actual vs Predicted Flight Delay')
plt.show()
kmeans = KMeans(n_clusters=3, random_state=42)
```

```
data['Cluster'] = kmeans.fit_predict(data[['Delay', 'Distance']])

plt.figure(figsize=(8, 6))

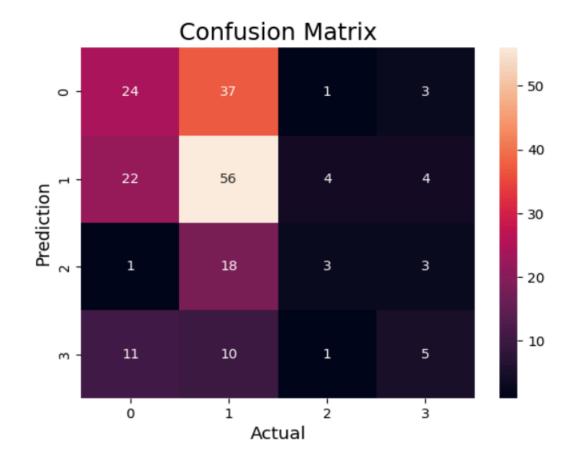
sns.scatterplot(x='Distance', y='Delay', hue='Cluster', data=data, palette='Set1')

plt.xlabel('Distance')

plt.ylabel('Delay')

plt.title('Clustering of Flights based on Delay and Distance')

plt.show()
```



| Classific | ation | Report: precision | recall | f1-score | support | |
|-----------|-------|----------------------|--------|----------|---------|--|
| | 0 | 0.41 | 0.37 | 0.39 | 65 | |
| | 1 | 0.46 | 0.65 | 0.54 | 86 | |
| | 2 | 0.33 | 0.12 | 0.18 | 25 | |
| | 3 | 0.33 | 0.19 | 0.24 | 27 | |
| accur | racy | | | 0.43 | 203 | |
| macro | avg | 0.39 | 0.33 | 0.34 | 203 | |
| weighted | avg | 0.41 | 0.43 | 0.41 | 203 | |
| | | | | | | |

RESULT:

Thus the program for predicting flight delays based on historical data is successfully executed and output is obtained.

| EX.N0:13 | TITANIC SURVIVAL PREDICTION |
|----------|-----------------------------|
| DATE: | |

To predict whether passengers survived the Titanic disaster.

ALGORITHM:

Step 1: Start the program.

Step 2: Import necessary libraries.

Step 3: Load the dataset.

Step 4: Extract and define feature in testing set and training set.

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

Step 6: Print equal metric & test the cell.

SOURCE CODE:

import pandas as pd

import numpy as np

import seaborn as sns

import matplotlib.pyplot as plt

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

```
titanic data = pd.read csv('tested.csv')
print("Missing values before handling:\n", titanic data.isnull().sum())
titanic data = titanic data.drop(['PassengerId', 'Name', 'Ticket', 'Cabin'], axis=1)
titanic data['Age'].fillna(titanic data['Age'].median(), inplace=True)
titanic data['Embarked'].fillna(titanic data['Embarked'].mode()[0], inplace=True)
titanic data['Fare'].fillna(titanic data['Fare'].median(), inplace=True)
print("Missing values after handling:\n", titanic data.isnull().sum())
titanic data = pd.get dummies(titanic data, columns=['Sex', 'Embarked'], drop first=True)
print(titanic data.head())
sns.countplot(x='Survived', data=titanic data)
plt.title('Survival Count')
plt.show()
plt.hist(titanic data['Age'], bins=20, color='lightblue')
plt.title('Age Distribution')
plt.xlabel('Age')
plt.ylabel('Frequency')
plt.show()
X = titanic data.drop('Survived', axis=1)
y = titanic data['Survived']
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_test = scaler.transform(X_test)

logreg = LogisticRegression()

logreg.fit(X_train, y_train)

y_pred = logreg.predict(X_test)

print("Accuracy:", accuracy_score(y_test, y_pred))

print("Confusion Matrix:\n", confusion_matrix(y_test, y_pred))

print("Classification Report:\n", classification report(y test, y pred))
```

| Confusion Matrix: [[50 0] [0 34]] Classification Report: | | | | | |
|---|-----------|--------|----------|---------|--|
| | precision | recall | f1-score | support | |
| 0 | 1.00 | 1.00 | 1.00 | 50 | |
| 1 | 1.00 | 1.00 | 1.00 | 34 | |
| accuracy | | | 1.00 | 84 | |
| macro avg | 1.00 | 1.00 | 1.00 | 84 | |
| weighted avg | 1.00 | 1.00 | 1.00 | 84 | |

PCA-Based Logistic Regression Accuracy: 0.6190476190476191

RESULT:

Thus the program to predict whether passengers survived the Titanic disaster is successfully executed and output is obtained.

| EX.N0 : 14 | ENERGY CONSUMPTION FORECASTING |
|------------|--------------------------------|
| DATE: | |

To forecast energy consumption based on historical data.

ALGORITHM:

Step 1: Start the program.

Step 2: Import necessary libraries.

Step 3: Load the dataset.

Step 4: Extract and define feature in testing set and training set.

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

Step 6: Print equal metric & test the cell.

SOURCE CODE:

import pandas as pd

import numpy as np

import seaborn as sns

import matplotlib.pyplot as plt

from sklearn.model selection import train test split

from sklearn.linear model import LinearRegression

from sklearn.metrics import mean squared error, accuracy score

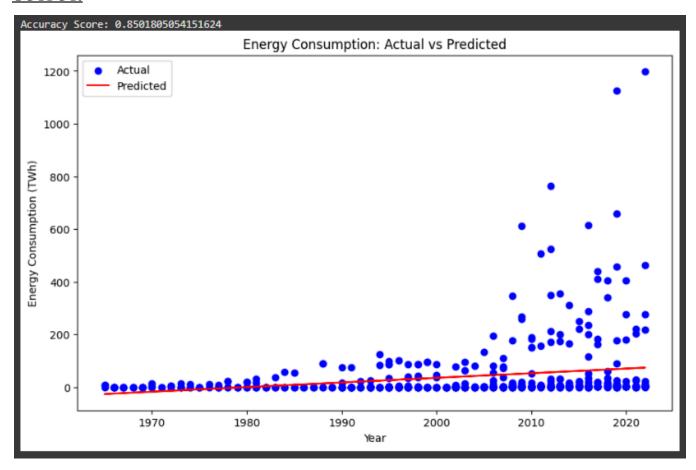
energy data = pd.read csv("World Energy Consumption.csv")

energy data clean = energy data[['year', 'biofuel consumption']].dropna()

print(energy data clean.head())

```
plt.figure(figsize=(10, 6))
plt.plot(energy data clean['year'], energy data clean['biofuel consumption'], marker='o')
plt.title('Energy Consumption Over Time')
plt.xlabel('Year')
plt.ylabel('Energy Consumption (TWh)')
plt.grid(True)
plt.show()
plt.figure(figsize=(10, 6))
sns.heatmap(energy data clean.corr(), annot=True, cmap='coolwarm')
plt.title('Correlation Heatmap of Energy Data')
plt.show()
X = energy_data_clean[['year']]
y = energy_data_clean['biofuel_consumption']
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
threshold = y_train.mean()
y_train_class = (y_train > threshold).astype(int)
y_test_class = (y_test > threshold).astype(int)
model = LinearRegression()
model.fit(X_train, y_train_class)
y_pred_class = (model.predict(X_test) > 0.5).astype(int)
accuracy = accuracy score(y test class, y pred class)
print(f"Accuracy Score: {accuracy}")
plt.figure(figsize=(10, 6))
plt.scatter(X test, y test, color='blue', label='Actual')
```

```
plt.plot(X_test, model.predict(X_test), color='red', label='Predicted')
plt.title('Energy Consumption: Actual vs Predicted')
plt.xlabel('Year')
plt.ylabel('Energy Consumption (TWh)')
plt.legend()
plt.show()
future_years = pd.DataFrame({'year': np.arange(2025, 2036)})
future_predictions = model.predict(future_years)
for year, consumption in zip(future_years['year'], future_predictions):
    print(f"Year {year}: Predicted Energy Consumption = {consumption:.2f} TWh")
```



```
Year 2025: Predicted Energy Consumption = 78.87 TWh
Year 2026: Predicted Energy Consumption = 80.63 TWh
Year 2027: Predicted Energy Consumption = 82.39 TWh
Year 2028: Predicted Energy Consumption = 84.15 TWh
Year 2029: Predicted Energy Consumption = 85.90 TWh
Year 2030: Predicted Energy Consumption = 87.66 TWh
Year 2031: Predicted Energy Consumption = 89.42 TWh
Year 2032: Predicted Energy Consumption = 91.18 TWh
Year 2033: Predicted Energy Consumption = 92.93 TWh
Year 2034: Predicted Energy Consumption = 94.69 TWh
Year 2035: Predicted Energy Consumption = 96.45 TWh
```

RESULT:

Thus the program to forecast energy consumption based on historical data is successfully executed and output is obtained.

EX.N0: 15 HUMAN ACTIVITY RECOGNITION DATE:

AIM:

To classify different human activities using smartphone sensor data.

ALGORITHM:

Step 1: Start the program.

Step 2: Import necessary libraries.

Step 3: Load the dataset.

Step 4: Extract and define feature in testing set and training set.

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

Step 6: Print equal metric & test the cell.

SOURCE CODE:

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.preprocessing import StandardScaler, LabelEncoder

from sklearn.decomposition import PCA

from sklearn.discriminant analysis import Linear Discriminant Analysis

from sklearn.ensemble import RandomForestClassifier

from sklearn.metrics import accuracy score, confusion matrix, classification report

```
data = pd.read csv("train.csv")
label encoder = LabelEncoder()
data['Activity'] = label encoder.fit transform(data['Activity'])
feature subset = data[['tBodyAcc-mean()-X', 'tBodyAcc-mean()-Y', 'tBodyAcc-mean()-Z',
'Activity']]
sns.pairplot(feature subset, hue="Activity", markers=".")
plt.show()
X = data.drop('Activity', axis=1)
y = data['Activity']
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})
pca = PCA(n components=2)
X train pca = pca.fit transform(X train)
X \text{ test pca} = \text{pca.transform}(X \text{ test})
plt.figure(figsize=(10, 7))
plt.scatter(X_train_pca[:, 0], X_train_pca[:, 1], c=y_train, cmap='rainbow', s=50)
plt.title('PCA of Training Data')
plt.xlabel('Principal Component 1')
plt.ylabel('Principal Component 2')
plt.show()
lda = LinearDiscriminantAnalysis(n components=2)
X train lda = lda.fit transform(X train, y train)
```

```
X test Ida = Ida.transform(X test)
plt.figure(figsize=(10, 7))
plt.scatter(X train lda[:, 0], X train lda[:, 1], c=y train, cmap='rainbow', s=50)
plt.title('LDA of Training Data')
plt.xlabel('Linear Discriminant 1')
plt.ylabel('Linear Discriminant 2')
plt.show()
clf = RandomForestClassifier(n estimators=100, random state=42)
clf.fit(X train, y train)
y pred = clf.predict(X test)
accuracy = accuracy score(y test, y pred)
print(f"Accuracy: {accuracy * 100:.2f}%")
print("Classification Report:\n", classification report(y test, y pred))
cm = confusion matrix(y test, y pred)
plt.figure(figsize=(10, 7))
sns.heatmap(cm, annot=True, fmt='d', cmap='Blues', xticklabels=label_encoder.classes_,
yticklabels=label encoder.classes )
plt.xlabel('Predicted')
plt.ylabel('True')
plt.title('Confusion Matrix')
plt.show()
```



| Accuracy: 97.96% | | | | | |
|------------------|-----------|--------|----------|---------|--|
| Classification | n Report: | | | | |
| | precision | recall | f1-score | support | |
| | | | | | |
| 0 | 1.00 | 1.00 | 1.00 | 280 | |
| 1 | 0.97 | 0.95 | 0.96 | 262 | |
| 2 | 0.96 | 0.97 | 0.96 | 276 | |
| 3 | 1.00 | 0.99 | 0.99 | 247 | |
| 4 | 0.99 | 0.98 | 0.98 | 206 | |
| 5 | 0.97 | 0.99 | 0.98 | 200 | |
| | | | | | |
| accuracy | | | 0.98 | 1471 | |
| macro avg | 0.98 | 0.98 | 0.98 | 1471 | |
| weighted avg | 0.98 | 0.98 | 0.98 | 1471 | |

RESULT:

Thus the program to classify different human activities using smartphone sensor data.