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.

VISUALIZATION: 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:

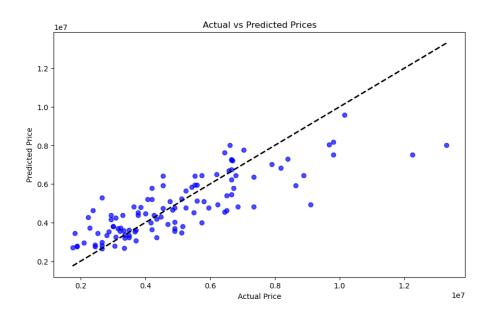
import pandas as pd

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/HARISH/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()
COMPUTATIONAL
                                                                                       221501034
```

```
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

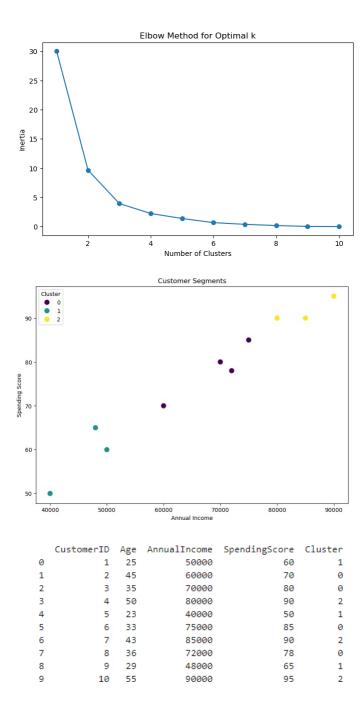
from sklearn.cluster import KMeans

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.

<u>VISUALIZATION:</u> 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

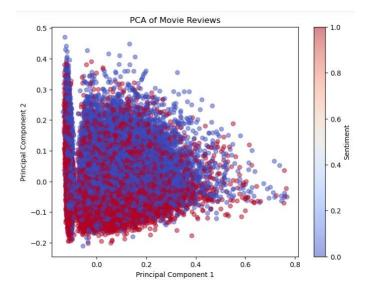
from wordcloud import WordCloud

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/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()
```



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

Classification	n 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.

VISUALIZATION: 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.

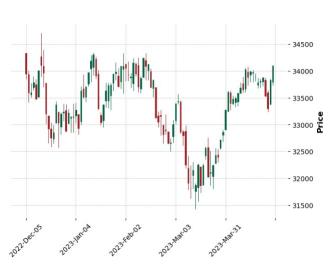
PROGRAM:

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

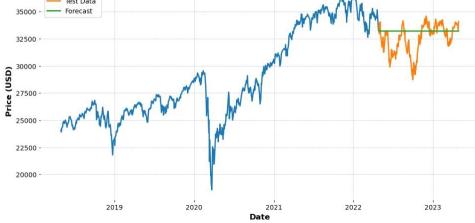
```
file path = r'C:\Users\HARISH\Downloads\yahoo data.xlsx'
data = pd.read excel(file path, index col='Date', parse dates=True)
data.rename(columns={'Close*': 'Close', 'Adj Close**': 'Adj Close'}, inplace=True)
data.sort index(inplace=True)
data.ffill(inplace=True)
if 'Adj Close' in data.columns:
plt.figure(figsize=(12, 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()
```

RMSE: 1517.2016308018185









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.

VISUALIZATION: 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

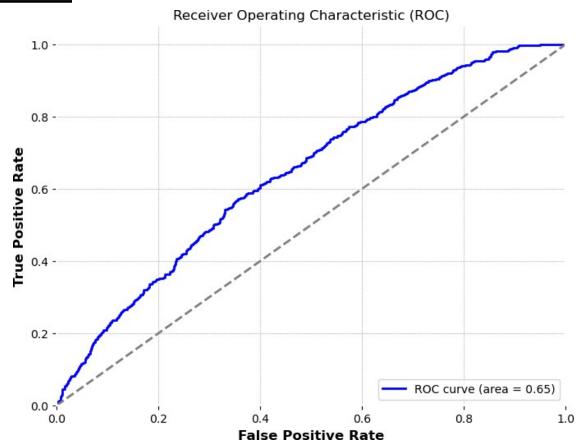
from sklearn.preprocessing import StandardScaler

from sklearn.decomposition import PCA

import os

```
file path = 'C:/Users/HARISH/Downloads/loan data.csv' # Update path
accordinglyif 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.

VISUALIZATION: 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

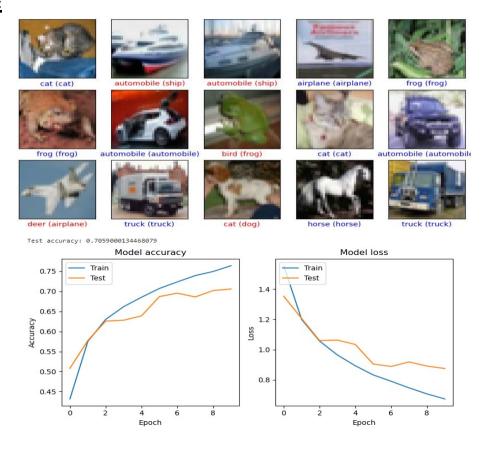
from tensorflow.keras.preprocessing.image import ImageDataGenerator

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	FREDICTING DIADETES

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

PYTHON CONCEPTS: Data structures, numeric types, functions.

VISUALIZATION: 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

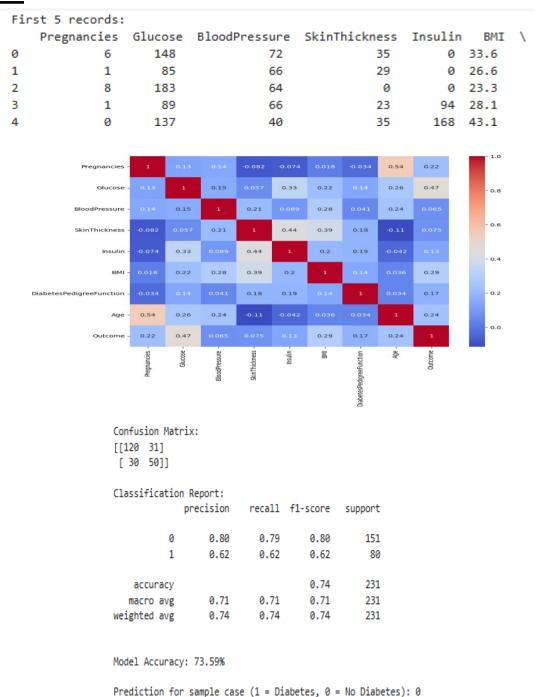
from sklearn.linear model import LogisticRegression

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]}")
```



RESULT:

Thus, the program for predicting diabetes is executed successfully.

8. WINE QUALITY 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.

<u>VISUALIZATION:</u> 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

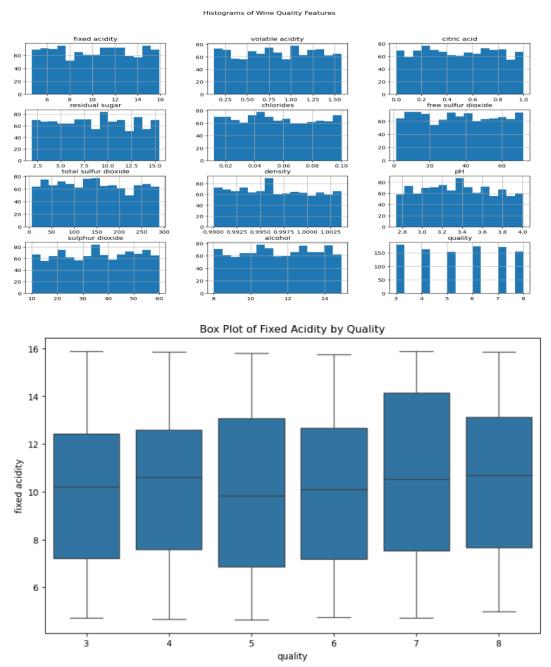
import seaborn as sns

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}')
```



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.

VISUALIZATION: 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

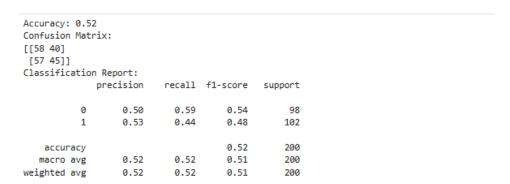
import seaborn as sns

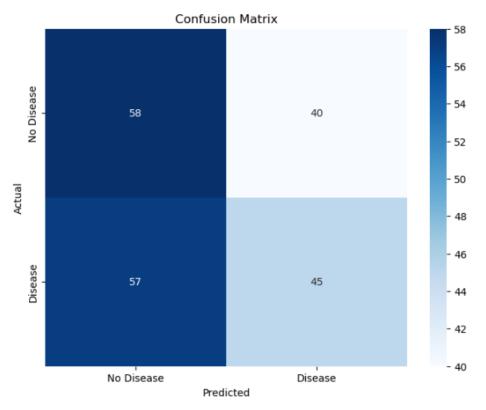
from sklearn.model selection import train test split

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) # For reproducibility
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()
```





RESULT:

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

10. BREAST CANCER DIAGNOSIS

EX.N0: 10	Procest Consequences
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

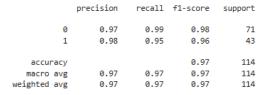
import seaborn as sns

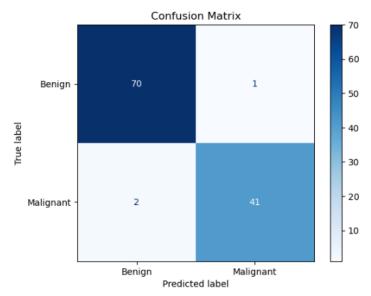
from sklearn.model_selection import train_test_split

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) # For reproducibility
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 area_se: 40.0 The tumor is predicted to be: Malignant

Based on the symptoms provided, the person may be at risk.

RESULT:

smoothness_se: 0.007

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

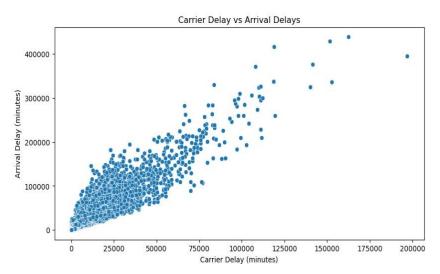
import seaborn as sns

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/HARISH/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
airport_name
                      0
arr_flights
                    240
                           8000
arr_del15
                    443
                    240
carrier_ct
                    240
weather_ct
nas_ct
                    240
security_ct
                    240
late_aircraft_ct
                    240
arr_cancelled
                    240
arr_diverted
                    240
                           2000
arr_delay
                    240
carrier_delay
                    240
weather_delay
                    240
nas_delay
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 =

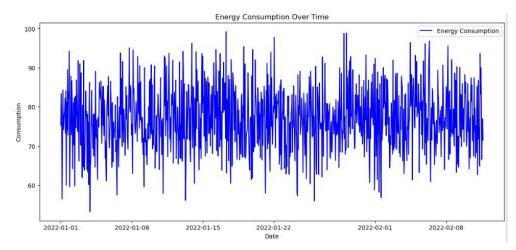
pd.read_csv('C:/Users/HARISH/Downloads/energy_consumption_dataset.csv',

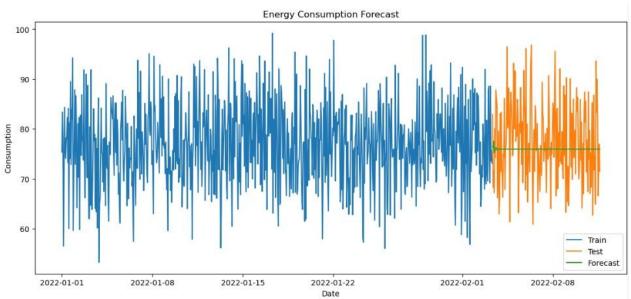
parse dates=['Timestamp'], index col='Timestamp')

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	\			HVACUsage Li	ghtingUsage	RenewableEnergy	DayOfWeek
Timestamp						Timestamp					
2022-01-01 00:00:00	25.139433	43.431581	1565.693999	5		2022-01-01	00:00:00	On	0ff	2.774699	Monday
2022-01-01 01:00:00	27.731651	54.225919	1411.064918	1		2022-01-01	01:00:00	On	On	21.831384	Saturday
2022-01-01 02:00:00	28.704277	58.907658	1755.715009	2		2022-01-01	02:00:00	0ff	0ff	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	0ff	3.071969	Friday





RESULT:

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