

**VISVESVARAYA TECHNOLOGICAL
UNIVERSITY**
“JnanaSangama”, Belgaum -590014, Karnataka.



**LAB REPORT
on**

Machine Learning (23CS6PCMAL)

Submitted by

S Gajana Nayak (1BM22CS227)

in partial fulfillment for the award of the degree of

**BACHELOR OF ENGINEERING
in
COMPUTER SCIENCE AND ENGINEERING**



**B.M.S. COLLEGE OF ENGINEERING
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B.M.S. College of Engineering,
Bull Temple Road, Bangalore 560019
(Affiliated To Visvesvaraya Technological University, Belgaum)

Department of Computer Science and Engineering



CERTIFICATE

This is to certify that the Lab work entitled “Machine Learning (23CS6PCMAL)” carried out by **S Gajanana Nayak (1BM22CS227)**, who is bonafide student of **B.M.S. College of Engineering**. It is in partial fulfillment for the award of **Bachelor of Engineering in Computer Science and Engineering** of the Visvesvaraya Technological University, Belgaum. The Lab report has been approved as it satisfies the academic requirements in respect of an Machine Learning (23CS6PCMAL) work prescribed for the said degree.

Lab Faculty Incharge	
Name: Ms. Saritha A N Assistant Professor Department of CSE, BMSCE	Dr. Kavitha Sooda Professor & HOD Department of CSE, BMSCE

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Github Link:

<https://github.com/Gajana227/ML>

Program 1

Write a python program to import and export data using Pandas library functions

Screenshot:

LAB - 01 3/8/25
Different ways of importing datasets:
Method-1: Introduce values directly Dataframe
import pandas as pd
data = {
 "USN": ["IBM23CSH17", "IBM22CS215", "IBM22CS227", "IBM22CS220",
 "IBM22CS214"],
 "Name": ["Rohit", "Rohit", "Gajanan", "Revanth", "Raghavendra"],
 "Marks": [85, 90, 88, 88, 92]
}
df = pd.DataFrame(data)
print(df)

Output:
USN Name Marks
0 IBM23CSH17 Rohit 85
1 IBM22CS215 Rohit 90
2 IBM22CS227 Gajanan 88
3 IBM22CS220 Revanth 88
4 IBM22CS214 Raghavendra 92

Method-2: Importing dataset from sklearn.datasets
from sklearn.datasets import load_diabetes
import pandas as pd
diabetes = load_diabetes()
df = pd.DataFrame(diabetes.data, columns=diabetes.feature_names)
df['target'] = diabetes.target
print(df.head())

Output:
Age Sex ... target
0 0.058 0.030 ... 245.0
1 -0.001 -0.044 ... 305.0
2 0.085 0.050 ... 206.0
3 -0.089 -0.044 0.012 235.0
4 0.005 -0.044 0.003 206.0

Method 3: Importing datasets from specific .csv file

Import pandas as pd
 $df = pd.read_csv('content/Diabetes.csv')$
 $print(df.head())$

Output:

	Gender	Age	BMI	CLASS
0	F	50	24.0	N
1	M	40	23.0	N
2	F	50	24.0	N
3	F	50	24.0	N
4	M	33	21.0	N

(odd man out)

Step 1: Analysing the Dataset

```

import yfinance as yf
import pandas as pd
import matplotlib.pyplot as plt
from datetime import date
from datetime import timedelta
from dateutil.relativedelta import relativedelta
import numpy as np
import warnings
warnings.filterwarnings('ignore')

# Fetching Data
tickers = ["HDFCBANK.NS", "ICICIBANK.NS"]
start = "2024-01-01"
end = "2024-12-31"

data = yf.download(tickers, start=start, end=end)
group_by_ticker = data.groupby("ticker")
print("First 5 rows of the dataset:")
print(data.head())
print("Shape of dataset:")
print(data.shape)
print("Column names:")
print(data.columns)

# Data Cleaning
hdfc_data = data['HDFCBANK.NS']
print("Summary stats for HDFC Bank:")
hdfc.describe()

hdfc['Daily Return'] = hdfc['Close'].pct_change()
hdfc['Daily Return'] = hdfc['Close'].pct_change()

icici_data = data['ICICIBANK.NS']
print("Summary stats for ICICI Bank:")
icici.describe()

icici['Daily Return'] = icici['Close'].pct_change()
icici['Daily Return'] = icici['Close'].pct_change()

```

```
kotak_data = data['KOTAKBANK.NS']
print("Summary stats for KOTAK Bank:")
print(kotak_data.describe())
kotak_data['Daily Return'] = kotak_data['Close'].pct_change()
kotak_data['Daily Return'] = kotak_data['Close'].pct_change()

plt.figure(figsize=(12,6))
plt.subplot(2,1,1)
hdfc_data['Close'].plot(title="HDFC Bank - Closing Price")
plt.subplot(2,1,2)
hdfc_data['Daily Return'].plot(title="HDFC Bank - Daily Returns",
                                color='orange')

plt.tight_layout()
plt.show()
hdfc_data.to_csv('hdfc-stock-data.csv')
print("HDFC stock data saved to 'hdfc-stock-data.csv'")
```

10
10

date
3/3/25

Code:

```
import pandas as pd

data = {

    "USN": ["1BM23CS417", "1BM22CS215", "1BM22CS227", "1BM22CS220", "1BM22CS214"],

    "Name": ["Rohit", "Rahul", "Gajanana", "Revanth", "Raghavendra"],

    "Marks": [85, 90, 78, 88, 92]

}

df = pd.DataFrame(data)

print(df)
```

```
from sklearn.datasets import load_diabetes

import pandas as pd

diabetes = load_diabetes()

df = pd.DataFrame(diabetes.data, columns=diabetes.feature_names)

df['target'] = diabetes.target

print(df.head())
```

```
import pandas as pd

df = pd.read_csv('content/Diabetes.csv')

print(df.head())

import yfinance as yf

import pandas as pd

import matplotlib.pyplot as plt

tickers = ["HDFCBANK.NS", "ICICIBANK.NS", "KOTAKBANK.NS"]
```

```
data = yf.download(tickers, start="2024-01-01", end="2024-12-30",
group_by='ticker')

print("First 5 rows of the dataset:")

print(data.head())

print("\nShape of the dataset:")

print(data.shape)

print("\nColumn names:")

print(data.columns)
```

```
hdfc_data = data['HDFCBANK.NS']

print("\nSummary statistics for HDFC Bank:")

print(hdfc_data.describe())

hdfc_data['Daily Return'] = hdfc_data['Close'].pct_change()

hdfc_data['Daily Return'] = hdfc_data['Close'].pct_change()

icici_data = data['ICICIBANK.NS']

print("\nSummary statistics for ICICI Bank:")

print(icici_data.describe())

icici_data['Daily Return'] = icici_data['Close'].pct_change()

icici_data['Daily Return'] = icici_data['Close'].pct_change()

kotak_data = data['KOTAKBANK.NS']

print("\nSummary statistics for KOTAK Bank:")

print(kotak_data.describe())

kotak_data['Daily Return'] = kotak_data['Close'].pct_change()

kotak_data['Daily Return'] = kotak_data['Close'].pct_change()
```

```
plt.figure(figsize=(12, 6))

plt.subplot(2, 1, 1)

hdfc_data['Close'].plot(title="HDFC Bank - Closing Price")

plt.subplot(2, 1, 2)

hdfc_data['Daily Return'].plot(title="HDFC Bank - Daily Returns", color='orange')

plt.tight_layout()

plt.show()

hdfc_data.to_csv('hdfc_stock_data.csv')

print("HDFC stock data saved to 'hdfc_stock_data.csv'.")


```

```
plt.figure(figsize=(12, 6))

plt.subplot(2, 1, 1)

icici_data['Close'].plot(title="ICICI Bank - Closing Price")

plt.subplot(2, 1, 2)

icici_data['Daily Return'].plot(title="ICICI Bank - Daily Returns", color='red')

plt.tight_layout()

plt.show()

icici_data.to_csv('icici_stock_data.csv')

print("ICICI stock data saved to 'icici_stock_data.csv'.")


```

```
plt.figure(figsize=(12, 6))

plt.subplot(2, 1, 1)

kotak_data['Close'].plot(title="Kotak Bank - Closing Price")
```

```

plt.subplot(2, 1, 2)

kotak_data['Daily Return'].plot(title="Kotak Bank - Daily Returns", color='green')

plt.tight_layout()

plt.show()

kotak_data.to_csv('kotak_stock_data.csv')

print("Kotak stock data saved to 'kotak_stock_data.csv'.")

```

Output:

	USN	Name	Marks
0	1BM23CS417	Rohit	85
1	1BM22CS215	Rahul	90
2	1BM22CS227	Gajana	78
3	1BM22CS220	Revanth	88
4	1BM22CS214	Raghavendra	92

	age	sex	bmi	bp	s1	s2	s3	\
0	0.038076	0.050680	0.061696	0.021872	-0.044223	-0.034821	-0.043401	
1	-0.001882	-0.044642	-0.051474	-0.026328	-0.008449	-0.019163	0.074412	
2	0.085299	0.050680	0.044451	-0.005670	-0.045599	-0.034194	-0.032356	
3	-0.089063	-0.044642	-0.011595	-0.036656	0.012191	0.024991	-0.036038	
4	0.005383	-0.044642	-0.036385	0.021872	0.003935	0.015596	0.008142	

	s4	s5	s6	target
0	-0.002592	0.019907	-0.017646	151.0
1	-0.039493	-0.068332	-0.092204	75.0
2	-0.002592	0.002861	-0.025930	141.0
3	0.034309	0.022688	-0.009362	206.0
4	-0.002592	-0.031988	-0.046641	135.0

	ID	No_Pation	Gender	AGE	Urea	Cr	HbA1c	Chol	TG	HDL	LDL	VLDL	\
0	502	17975	F	50	4.7	46	4.9	4.2	0.9	2.4	1.4	0.5	
1	735	34221	M	26	4.5	62	4.9	3.7	1.4	1.1	2.1	0.6	
2	420	47975	F	50	4.7	46	4.9	4.2	0.9	2.4	1.4	0.5	
3	680	87656	F	50	4.7	46	4.9	4.2	0.9	2.4	1.4	0.5	
4	504	34223	M	33	7.1	46	4.9	4.9	1.0	0.8	2.0	0.4	

	BMI	CLASS
0	24.0	N
1	23.0	N
2	24.0	N
3	24.0	N
4	21.0	N

First 5 rows of the dataset:

Ticker	HDFCBANK.NS				
Price	Open	High	Low	Close	Volume
Date					
2024-01-01	1683.017598	1686.125187	1669.206199	1675.223999	7119843
2024-01-02	1675.914685	1679.860799	1665.950651	1676.210571	14621046
2024-01-03	1679.071480	1681.735059	1646.466666	1650.363525	14194881
2024-01-04	1655.394910	1672.116520	1648.193203	1668.071777	13367028
2024-01-05	1664.421596	1681.932477	1645.628180	1659.538208	15944735

Ticker	KOTAKBANK.NS				
Price	Open	High	Low	Close	Volume
Date					
2024-01-01	1906.909954	1916.899006	1891.027338	1907.059814	1425902
2024-01-02	1905.911108	1905.911108	1858.063525	1863.008179	5120796
2024-01-03	1861.959234	1867.952665	1845.627158	1863.857178	3781515
2024-01-04	1869.451068	1869.451068	1858.513105	1861.559692	2865766
2024-01-05	1863.457575	1867.852782	1839.383985	1845.577148	7799341

Ticker	ICICIBANK.NS				
Price	Open	High	Low	Close	Volume
Date					
2024-01-01	983.086778	996.273246	982.541485	990.869812	7683792
2024-01-02	988.490253	989.134730	971.883221	973.866150	16263825
2024-01-03	976.295294	979.567116	966.777197	975.650818	16826752
2024-01-04	977.980767	980.707295	973.519176	978.724365	22789140
2024-01-05	979.567084	989.779158	975.402920	985.218445	14875499

Shape of the dataset:
(244, 15)

Column names:

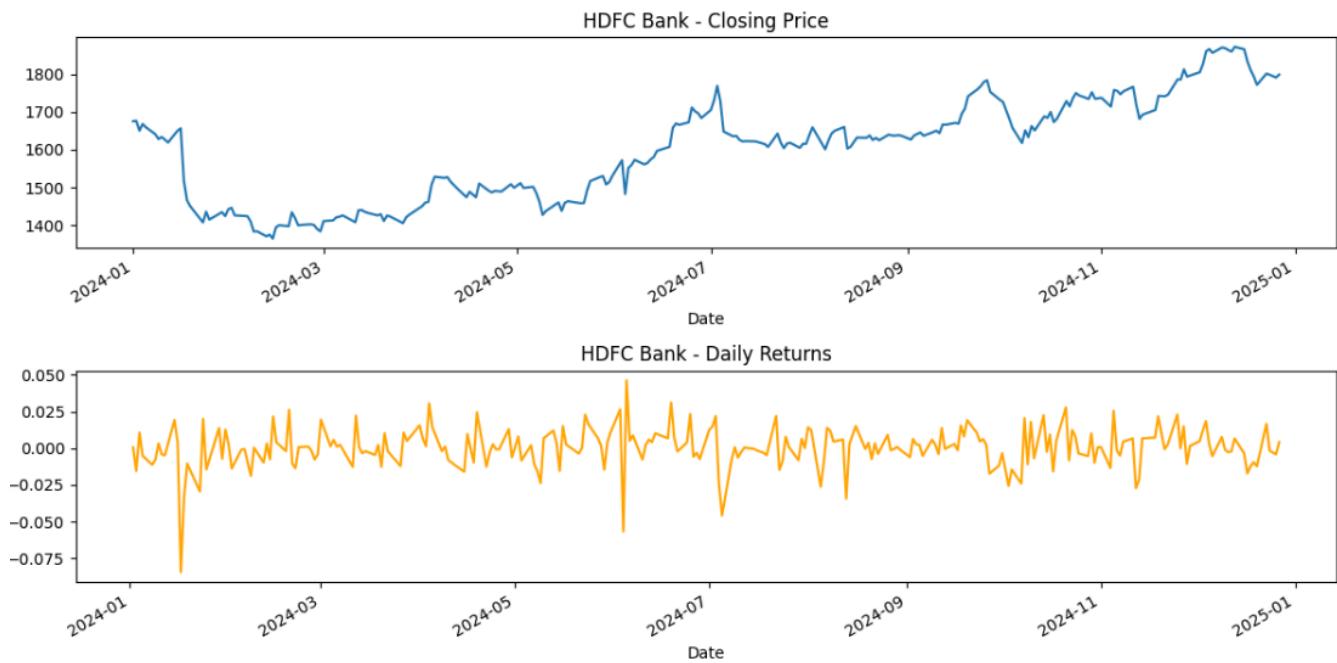
```
MultiIndex([('HDFCBANK.NS', 'Open'),
            ('HDFCBANK.NS', 'High'),
            ('HDFCBANK.NS', 'Low'),
            ('HDFCBANK.NS', 'Close'),
            ('HDFCBANK.NS', 'Volume'),
            ('KOTAKBANK.NS', 'Open'),
            ('KOTAKBANK.NS', 'High'),
            ('KOTAKBANK.NS', 'Low'),
            ('KOTAKBANK.NS', 'Close'),
            ('KOTAKBANK.NS', 'Volume'),
            ('ICICIBANK.NS', 'Open'),
            ('ICICIBANK.NS', 'High'),
            ('ICICIBANK.NS', 'Low'),
            ('ICICIBANK.NS', 'Close'),
            ('ICICIBANK.NS', 'Volume')],
           names=['Ticker', 'Price'])
```

Summary statistics for HDFC Bank:

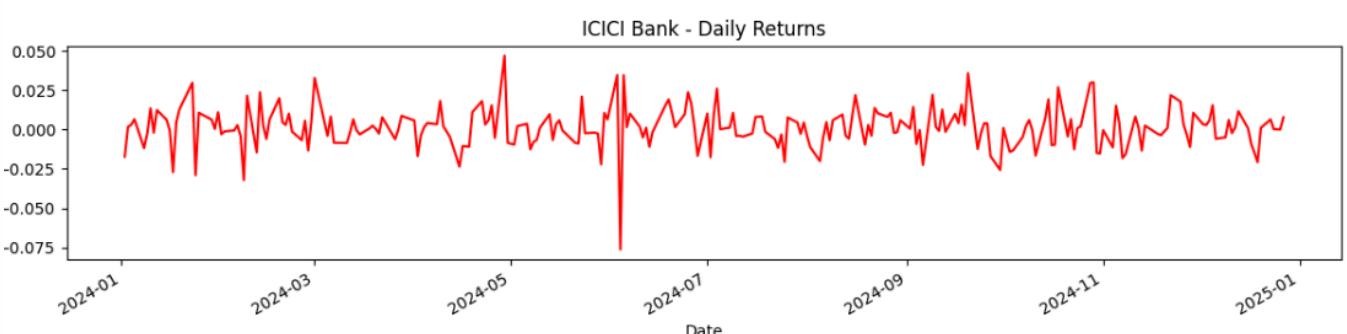
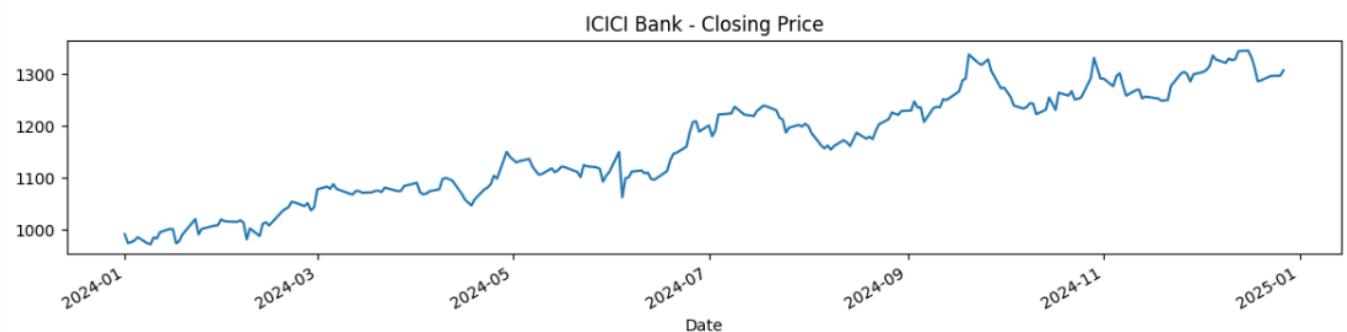
Price	Open	High	Low	Close	Volume
count	244.000000	244.000000	244.000000	244.000000	2.440000e+02
mean	1601.375295	1615.443664	1588.221245	1601.898968	2.119658e+07
std	134.648125	134.183203	132.796819	133.748372	2.133860e+07
min	1357.463183	1372.754374	1345.180951	1365.404785	8.798460e+05
25%	1475.316358	1494.072805	1460.259509	1474.564087	1.274850e+07
50%	1627.724976	1638.350037	1616.000000	1625.950012	1.686810e+07
75%	1696.474976	1711.425018	1679.250000	1697.062531	2.295014e+07
max	1877.699951	1880.000000	1858.550049	1871.750000	2.226710e+08

```
Summary statistics for ICICI Bank:
Price      Open      High       Low      Close      Volume
count    244.000000  244.000000  244.000000  244.000000  2.440000e+02
mean    1161.723560 1173.687900 1151.318979 1162.751791 1.539172e+07
std     104.905646 105.668229 105.083015 105.520481 9.503609e+06
min     965.637027 979.567116 961.869473 971.387512 1.007022e+06
25%    1073.818215 1085.368782 1067.386038 1075.107086 1.014533e+07
50%    1169.443635 1178.450012 1157.361521 1165.470703 1.291768e+07
75%    1248.512512 1261.399994 1236.649963 1250.812531 1.755770e+07
max    1344.900024 1362.349976 1340.050049 1346.099976 7.325777e+07
```

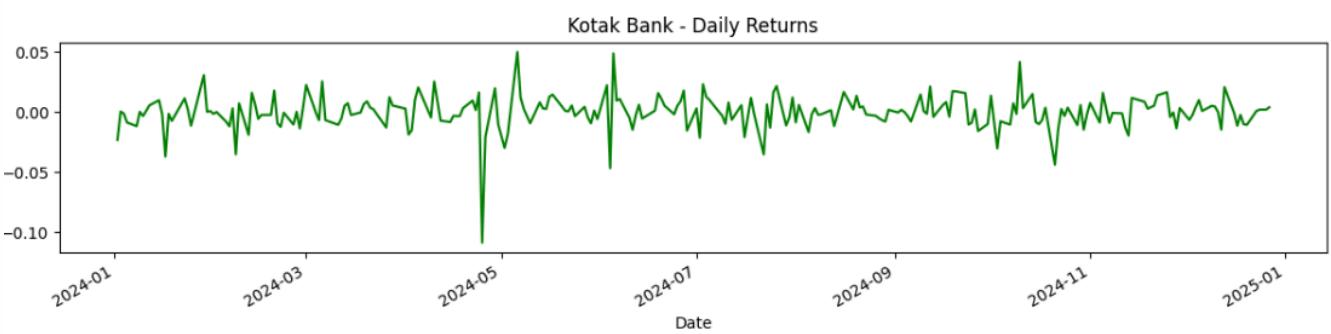
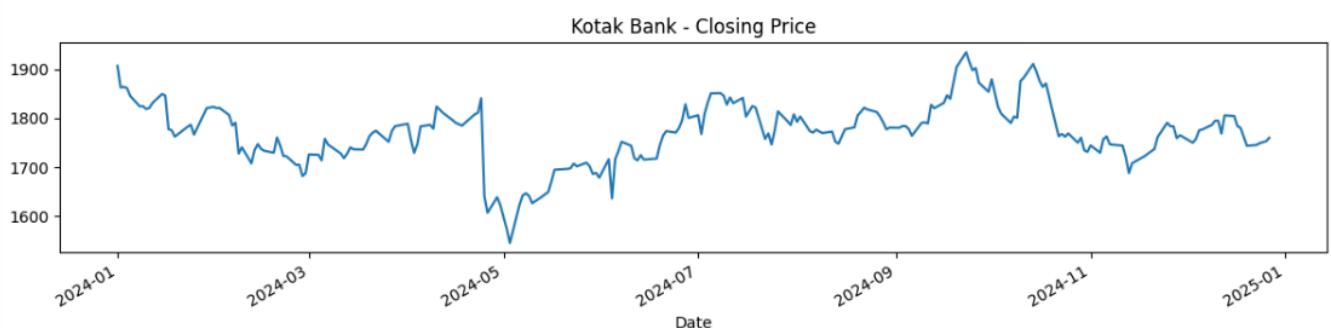
```
Summary statistics for KOTAK Bank:
Price      Open      High       Low      Close      Volume
count    244.000000  244.000000  244.000000  244.000000  2.440000e+02
mean    1771.245907 1787.548029 1754.395105 1770.792347 5.736598e+06
std     62.189675 61.978802 62.765980 62.594747 5.388927e+06
min     1581.266899 1586.161558 1542.159736 1545.006592 1.824890e+05
25%    1733.974927 1754.131905 1719.028421 1736.297058 3.300380e+06
50%    1769.500000 1789.450012 1758.099976 1773.681030 4.307680e+06
75%    1809.925018 1826.998164 1789.912506 1808.155670 6.159475e+06
max    1935.000000 1942.000000 1909.599976 1934.699951 6.617908e+07
```



```
HDFC stock data saved to 'hdfc_stock_data.csv'.
```



ICICI stock data saved to 'icici_stock_data.csv'.



Kotak stock data saved to 'kotak_stock_data.csv'.

Program 2

Demonstrate various data pre-processing techniques for a given dataset

Screenshot:

LAB-1 :

```

write python code, consider filename as "housing.csv"
    i) To load .csv file into the dataframe
        import pandas as pd
        filename = "housing.csv"
        df = pd.read_csv(filename)
    ii) To display information of all columns
        print(df.info())
    iii) To display statistical information of all numerical
        print(df.describe())
    iv) To display the count of unique labels for "Ocean Proximity" column
        if "Ocean Proximity" in df.columns:
            print(df["Ocean Proximity"].value_counts())
        else:
            print("Not found")
    v) To display which attributes in dataset have missing values
        count greater than zero
        m_values = df.isnull().sum()
        m_columns = m_values[m_values > 0]
        if not m_columns.empty:
            print(m_columns)
        else:
            print("No missing values found")

```

Output:

Index	Attribute	Count	Non-null	Data Type
0	longitude	3000	non-null	float64
1	latitude	3000	non-null	float64
2	housing_median_age	3000	non-null	float64
3	total_rooms	3000	non-null	float64
4	total_bedrooms	3000	non-null	float64
5	population	3000	non-null	float64
6	households	3000	non-null	float64
7	median_income	3000	non-null	float64
8	median_house_value	3000	non-null	float64

v) Longitude, Latitude and Wind direction variables are in column 23
 wind speed and wind direction are present in row 23

	Longitude	Latitude	Wind Speed	Wind Direction
count	3000	3000	3000	3000
mean	-119.5	35.6	3.19	210
std	1.99	3.05	3.05	3.05
min	-124.18	33.9	0	0
25%	-121.81	34.2	0	0
50%	-118.48	34.2	0	0
75%	-118.02	37.6	240	240
max	-114.19	41.9	240	240

vi) 'Ocean Proximity' column not found in the dataset

vii) No missing values found in the dataset

Question: several 204 bio books had missing volumes. How did columns in the database had missing values?

Q) Which columns in the dataset had missing values? How do you handle them?

get: Missing values columns:
Adult Income dataset → Age, Salary
Iris dataset → Glucose, BMI

Diabetes dataset → glucose / ...
using approach:
... random since it's lost

Handling approach: \rightarrow ~~1st~~ → 2nd egg \rightarrow used medium since its last
survivors

Handling outliers → Income distribution → for e.g. → used Median
sensitive to outliers

Diabetes dataset → Glucose → used median since glucose levels may have outliers
→ BMI → used mean assuming normal distribution.

Q2) Which categorical columns did you identify in the dataset?
How did you encode them?

80%: Adult Income Deficit

80%: Adult Income Dataset:
Categorical columns: Gender → original Encoding
City → One-Hot Encoding

Diabetes Dosages

Diabetes Dataset :
categorical columns : Gender → original Encoding
Outcome → One-Hot Encoding.

Q3) what is the difference between Min-Max scaling and Standardization? when would you use one over the other?

Soln: Min-Max Scaling:

$$\rightarrow X' = \frac{X - X_{\min}}{X_{\max} - X_{\min}}$$

→ scales values between 0 & 1.

→ less affected by outliers

→ sensitive to outliers

Standardization

$$\rightarrow X' = \frac{X - \mu}{\sigma}$$

→ transforms data to have mean = 0 and variance = 1

when data is not normally distributed and has known bounds, min-max scaling is used.

when data follows a normal distribution, standardization is used.

When dealing with outliers, standardization is preferred because it is less affected by outliers.

Normalizing some features helps in standardizing the features.

Linear regression needs both standardization and normalization.

If standardization is not applied, then the regression coefficients will be very large.

standardization is better than normalization.

standardization is better than normalization.

Code:

```
import pandas as pd
```

```
import numpy as np
```

```
import matplotlib.pyplot as plt
```

```

from sklearn.model_selection import train_test_split

from sklearn.impute import SimpleImputer

from sklearn.preprocessing import OrdinalEncoder, OneHotEncoder

from sklearn.preprocessing import StandardScaler, MinMaxScaler

from scipy import stats

filename = "/content/housing.csv"

df = pd.read_csv(filename)

print("Dataset Information:")

print(df.info())

print("\nStatistical Summary of Numerical Columns:")

print(df.describe())


if "Ocean Proximity" in df.columns:

    print("\nUnique Value Counts for 'Ocean Proximity':")

    print(df["Ocean Proximity"].value_counts())

else:

    print("\n'Ocean Proximity' column not found in the dataset.")


missing_values = df.isnull().sum()

missing_columns = missing_values[missing_values > 0]


if not missing_columns.empty:

    print("\nColumns with Missing Values:")

    print(missing_columns)

```

```
else:  
    print("\nNo missing values found in the dataset.")  
  
'''Write Python code to implement the following data preprocessing  
techniques for Adult income data set'''
```

```
def createdata():  
  
    data = {  
  
        'Age': np.random.randint(18, 70, size=20),  
  
        'Salary': np.random.randint(30000, 120000, size=20),  
  
        'Purchased': np.random.choice([0, 1], size=20),  
  
        'Gender': np.random.choice(['Male', 'Female'], size=20),  
  
        'City': np.random.choice(['New York', 'San Francisco', 'Los Angeles'], size=20)  
  
    }  
  
    df = pd.DataFrame(data)  
  
    return df  
  
df = createdata()  
  
df.head(10)  
  
# Introduce some missing values for demonstration  
  
df.loc[5, 'Age'] = np.nan  
  
df.loc[10, 'Salary'] = np.nan  
  
df.head(10)
```

```

#Data Cleaning

# Check for missing values in each column

missing_values = df.isnull().sum()

# Display columns with missing values

print(missing_values[missing_values > 0])

#Set the values to some value (zero, the mean, the median, etc.).

# Step 1: Create an instance of SimpleImputer with the median strategy for Age and mean stratergy for Salary

imputer1 = SimpleImputer(strategy="median")

imputer2 = SimpleImputer(strategy="mean")

df_copy=df

# Step 2: Fit the imputer on the "Age" and "Salary"column

# Note: SimpleImputer expects a 2D array, so we reshape the column

imputer1.fit(df_copy[["Age"]])

imputer2.fit(df_copy[["Salary"]])

# Step 3: Transform (fill) the missing values in the "Age" and "Salary"column

df_copy["Age"] = imputer1.transform(df[["Age"]])

df_copy["Salary"] = imputer2.transform(df[["Salary"]])

# Verify that there are no missing values left

print(df_copy["Age"].isnull().sum())

print(df_copy["Salary"].isnull().sum())

#Handling Categorical Attributes

#Using Ordinal Encoding for gender COlumn and One-Hot Encoding for City Column

# Initialize OrdinalEncoder

ordinal_encoder = OrdinalEncoder(categories=[["Male", "Female"]])

```

```

# Fit and transform the data

df_copy["Gender_Encoded"] = ordinal_encoder.fit_transform(df_copy[["Gender"]])

# Initialize OneHotEncoder

onehot_encoder = OneHotEncoder()

# Fit and transform the "City" column

encoded_data = onehot_encoder.fit_transform(df[["City"]])

# Convert the sparse matrix to a dense array

encoded_array = encoded_data.toarray()

# Convert to DataFrame for better visualization

encoded_df = pd.DataFrame(encoded_array,
columns=onehot_encoder.get_feature_names_out(["City"]))

df_encoded = pd.concat([df_copy, encoded_df], axis=1)

df_encoded.drop("Gender", axis=1, inplace=True)

df_encoded.drop("City", axis=1, inplace=True)

print(df_encoded. head())

```

#Removing Outliers

Outlier Detection and Treatment using IQR

#Pros: Simple and effective for mild outliers.

#Cons: May overly reduce variation if there are many extreme outliers.

```

df_encoded_copy1=df_encoded

df_encoded_copy2=df_encoded

df_encoded_copy3=df_encoded

```

Q1 = df_encoded_copy1['Salary'].quantile(0.25)

```
Q3 = df_encoded_copy1['Salary'].quantile(0.75)

IQR = Q3 - Q1

lower_bound = Q1 - 1.5 * IQR

upper_bound = Q3 + 1.5 * IQR

df_encoded_copy1['Salary'] = np.where(df_encoded_copy1['Salary'] > upper_bound, upper_bound,
                                       np.where(df_encoded_copy1['Salary'] < lower_bound, lower_bound,
                                                df_encoded_copy1['Salary']))

df_encoded_copy1.head()
```

#Data Transformation

Min-Max Scaler/Normalization (range 0-1)

#Pros: Keeps all data between 0 and 1; ideal for distance-based models.

#Cons: Can distort data distribution, especially with extreme outliers.

```
normalizer = MinMaxScaler()
```

```
df_encoded[['Salary']] = normalizer.fit_transform(df_encoded[['Salary']])
```

```
df_encoded.head()
```

Standardization (mean=0, variance=1)

#Pros: Works well for normally distributed data; suitable for many models.

#Cons: Sensitive to outliers.

```
scaler = StandardScaler()
```

```
df_encoded[['Age']] = scaler.fit_transform(df_encoded[['Age']])
```

```
df_encoded.head()
```

"Write Python code to implement the following data preprocessing

techniques for Diabetes data set"

```
def create_diabetes_data():
```

```
    data = {
```

```
        'Age': np.random.randint(20, 80, size=20),
```

```
        'Glucose': np.random.randint(50, 200, size=20),
```

```
        'BMI': np.random.uniform(18.5, 45.0, size=20),
```

```
        'Gender': np.random.choice(['Male', 'Female'], size=20),
```

```
        'Outcome': np.random.choice([0, 1], size=20)
```

```
}
```

```
df = pd.DataFrame(data)
```

Introduce some missing values for demonstration

```
df.loc[5, 'Glucose'] = np.nan
```

```
df.loc[10, 'BMI'] = np.nan
```

```
return df
```

Load dataset

```
df = create_diabetes_data()
```

```
print("Original Data:\n", df.head())
```

Data Cleaning - Handling Missing Values

```
imputer_glucose = SimpleImputer(strategy="median")
```

```
imputer_bmi = SimpleImputer(strategy="mean")
```

```
df_copy = df.copy()
```

```
df_copy["Glucose"] = imputer_glucose.fit_transform(df_copy[["Glucose"]])
```

```
df_copy["BMI"] = imputer_bmi.fit_transform(df_copy[["BMI"]])
```

```

print(df_copy["Glucose"].isnull().sum())

print(df_copy["BMI"].isnull().sum())

# Handling Categorical Attributes

# Ordinal Encoding for Gender

ordinal_encoder = OrdinalEncoder(categories=[["Male", "Female"]])

df_copy["Gender_Encoded"] = ordinal_encoder.fit_transform(df_copy[["Gender"]])

# One-Hot Encoding for Outcome column

onehot_encoder = OneHotEncoder()

encoded_outcome = onehot_encoder.fit_transform(df_copy[["Outcome"]]).toarray()

encoded_df = pd.DataFrame(encoded_outcome,
columns=onehot_encoder.get_feature_names_out(["Outcome"]))

df_encoded = pd.concat([df_copy, encoded_df], axis=1)

df_encoded.drop(["Gender", "Outcome"], axis=1, inplace=True)

print("\nAfter Encoding:\n", df_encoded.head())

```

```

# Removing Outliers using IQR (for Glucose)

Q1 = df_encoded["Glucose"].quantile(0.25)

Q3 = df_encoded["Glucose"].quantile(0.75)

IQR = Q3 - Q1

lower_bound = Q1 - 1.5 * IQR

upper_bound = Q3 + 1.5 * IQR

df_encoded["Glucose"] = np.where(df_encoded["Glucose"] > upper_bound, upper_bound,
np.where(df_encoded["Glucose"] < lower_bound, lower_bound,
df_encoded["Glucose"]))

```

```

df_encoded.head()

# Data Transformation

# Min-Max Scaling for BMI

normalizer = MinMaxScaler()

df_encoded[['BMI']] = normalizer.fit_transform(df_encoded[['BMI']])

df_encoded.head()

# Standardization for Age

scaler = StandardScaler()

df_encoded[['Age']] = scaler.fit_transform(df_encoded[['Age']])

print("\nFinal Preprocessed Data:\n", df_encoded.head())

```

Output:

```

Statistical Summary of Numerical Columns:
      longitude    latitude  housing_median_age  total_rooms \
count  3000.000000  3000.000000  3000.000000  3000.000000
mean   -119.589200    35.63539    28.845333  2599.578667
std     1.994936    2.12967    12.555396  2155.593332
min    -124.180000   32.56000    1.000000   6.000000
25%   -121.810000   33.93000    18.000000  1401.000000
50%   -118.485000   34.27000    29.000000  2106.000000
75%   -118.020000   37.69000    37.000000  3129.000000
max    -114.490000   41.92000    52.000000  30450.000000

      total_bedrooms  population  households  median_income \
count  3000.000000  3000.000000  3000.000000  3000.000000
mean   529.950667  1402.798667  489.91200   3.807272
std    415.654368  1030.543012  365.42271   1.854512
min     2.000000    5.000000    2.000000   0.499900
25%   291.000000   780.000000  273.000000   2.544000
50%   437.000000  1155.000000  409.50000   3.487150
75%   636.000000  1742.750000  597.25000   4.656475
max    5419.000000 11935.000000 4930.00000  15.000100

      median_house_value
count            3000.00000
mean            205846.27500
std             113119.68747
min            22500.00000
25%           121200.00000
50%           177650.00000
75%           263975.00000
max            500001.00000

'Ocean Proximity' column not found in the dataset.

No missing values found in the dataset.

```

	Age	Salary	Purchased	Gender	City
0	21	98335	0	Female	San Francisco
1	29	107193	1	Female	San Francisco
2	40	65298	0	Male	Los Angeles
3	67	110619	0	Female	New York
4	35	35539	1	Female	Los Angeles
5	52	119090	1	Male	New York
6	40	87266	0	Male	San Francisco
7	20	118367	1	Male	Los Angeles
8	48	102695	0	Female	San Francisco
9	26	66367	0	Male	San Francisco

```

    Age      Salary Purchased Gender_Encoded City_Los Angeles City_New York \
0 21.0    98335.0      0            1.0        0.0        0.0
1 29.0   107193.0      1            1.0        0.0        0.0
2 40.0    65298.0      0            0.0        1.0        0.0
3 67.0   110619.0      0            1.0        0.0        1.0
4 35.0    35539.0      1            1.0        1.0        0.0

City_San Francisco
0            1.0
1            1.0
2            0.0
3            0.0
4            0.0

```

	Age	Salary	Purchased	Gender_Encoded	City_Los Angeles	City_New York	City_San Francisco
0	21.0	0.745689	0	1.0	0.0	0.0	1.0
1	29.0	0.850876	1	1.0	0.0	0.0	1.0
2	40.0	0.353382	0	0.0	1.0	0.0	0.0
3	67.0	0.891559	0	1.0	0.0	1.0	0.0
4	35.0	0.000000	1	1.0	1.0	0.0	0.0

	Age	Glucose	BMI	Gender_Encoded	Outcome_0	Outcome_1
0	1.982577	93.0	0.991819	1.0	1.0	0.0
1	-0.110300	195.0	0.435823	0.0	0.0	1.0
2	0.229085	87.0	0.264387	1.0	0.0	1.0
3	-0.279993	113.0	0.785170	1.0	0.0	1.0
4	-1.015328	163.0	0.144165	0.0	0.0	1.0

```

Original Data:
   Age  Glucose      BMI  Gender  Outcome
0    77    93.0  44.640289  Female     0
1    40   195.0  30.589381   Male      1
2    46    87.0  26.256915  Female     1
3    37   113.0  39.417936  Female     1
4    24   163.0  23.218723   Male      1

```

	Age	Glucose	BMI	Gender_Encoder	Outcome_0	Outcome_1
0	77	93.0	0.991819	1.0	1.0	0.0
1	40	195.0	0.435823	0.0	0.0	1.0
2	46	87.0	0.264387	1.0	0.0	1.0
3	37	113.0	0.785170	1.0	0.0	1.0
4	24	163.0	0.144165	0.0	0.0	1.0

```

Final Preprocessed Data:
   Age  Glucose      BMI  Gender_Encoder  Outcome_0  Outcome_1
0  1.982577    93.0  0.991819        1.0       1.0       0.0
1 -0.110300   195.0  0.435823        0.0       0.0       1.0
2  0.229085    87.0  0.264387        1.0       0.0       1.0
3 -0.279993   113.0  0.785170        1.0       0.0       1.0
4 -1.015328   163.0  0.144165        0.0       0.0       1.0

```

Program 3

Implement Linear and Multi-Linear Regression algorithm using appropriate dataset

Screenshot:

LAB - 3
Simple Linear Regression

```

import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
# Data points
x = [10, 12, 14, 16, 18, 20]
y = [15, 17, 19, 21, 23, 25]

def estimate_coeff(x, y):
    n = np.size(x)
    m_x = np.mean(x)
    m_y = np.mean(y)
    ss_xy = np.sum((x - m_x) * (y - m_y))
    ss_xx = np.sum((x - m_x) ** 2)
    b_1 = ss_xy / ss_xx
    b_0 = m_y - b_1 * m_x
    return [b_0, b_1]

def plot(x, y, b):
    plt.scatter(x, y, color='m')
    y_pred = b[0] + b[1] * x
    plt.plot(x, y_pred, color='g')
    plt.xlabel('x')
    plt.ylabel('y')
    plt.show()

file_path = input("Enter path:")
df = pd.read_csv(file_path)

x = df.iloc[:, 0].values
y = df.iloc[:, 1].values
b = estimate(x, y)
plot(x, y, b)

```

Output:
 Enter path : /content/tumarketing.csv
 Estimated coefficients:
 $b_0 = 7.032$
 $b_1 = 0.047$

Multiple Linear Regression

```
data = [[1, 2, 3, 4, 5, 6, 7, 8, 9, 10],  
        [2, 3, 5, 7, 11, 13, 17, 19, 23, 29],  
        [3, 6, 9, 12, 15, 18, 21, 24, 27, 30],  
        [5, 9, 15, 22, 31, 41, 53, 61, 80, 96]]  
  
df = pd.DataFrame(data)  
X = df.drop(columns=["Target"], axis=1).values  
y = df["Target"].values.reshape(-1, 1)  
  
X = np.hstack((np.ones((X.shape[0], 1)), X))  
  
beta = np.linalg.solve(X.T @ X + 0.01 * np.identity(X.shape[1]), X.T @ y)  
  
y_pred = X @ beta  
  
mse = np.mean((y - y_pred) ** 2)  
tot_var = np.sum((y - np.mean(y)) ** 2)  
exp_var = np.sum((y_pred - np.mean(y)) ** 2)  
r2 = exp_var / tot_var  
  
print("Output:")  
  
Output:  
Model Coefficients: [0.040, 3.313, 0.120]  
Intercept: -3.1599  
Mean Squared Error: 5.362  
R-Squared Score: 0.993
```

Code:

```
#Linear Regression

import numpy as np

import pandas as pd

import matplotlib.pyplot as plt

def estimate_coef(x, y):

    n = np.size(x)

    m_x = np.mean(x)

    m_y = np.mean(y)

    SS_xy = np.sum((x - m_x) * (y - m_y))

    SS_xx = np.sum((x - m_x) ** 2)

    b_1 = SS_xy / SS_xx

    b_0 = m_y - b_1 * m_x

    return (b_0, b_1)

def plot_regression_line(x, y, b):

    plt.scatter(x, y, color="m", marker="o", s=30)

    y_pred = b[0] + b[1] * x

    plt.plot(x, y_pred, color="g")

    plt.xlabel('x')

    plt.ylabel('y')

    plt.title("Linear Regression")

    plt.show()

#Load dataset

file_path = input("Enter the path to the CSV file: ")
```

```

df = pd.read_csv(file_path)

# Assuming the dataset has two numerical columns: 'x' and 'y'

x = df.iloc[:, 0].values # First column as x

y = df.iloc[:, 1].values # Second column as y

b = estimate_coef(x, y)

print(f"Estimated coefficients:\nb_0 = {b[0]}\nb_1 = {b[1]}")

plot_regression_line(x, y, b)

```

#Multiple Regression

```

import numpy as np

import pandas as pd

import matplotlib.pyplot as plt

# Sample dataset

data = {
    "Feature1": [1, 2, 3, 4, 5, 6, 7, 8, 9, 10],
    "Feature2": [2, 3, 5, 7, 11, 13, 17, 19, 23, 29],
    "Feature3": [3, 6, 9, 12, 15, 18, 21, 24, 27, 30],
    "Target": [5, 9, 15, 22, 31, 41, 53, 66, 80, 96]
}

```

```
df = pd.DataFrame(data)
```

```

# Split dataset into features (X) and target variable (y)

X = df.drop(columns=["Target"]).values

y = df["Target"].values.reshape(-1, 1)

```

```

# Add intercept column (bias term)
X = np.hstack((np.ones((X.shape[0], 1)), X))

# Compute the coefficients using the Normal Equation
beta = np.linalg.solve(X.T @ X + 0.01 * np.identity(X.shape[1]), X.T @ y)

# Make predictions
y_pred = X @ beta

# Evaluate the model
mse = np.mean((y - y_pred) ** 2)
total_variance = np.sum((y - np.mean(y)) ** 2)
explained_variance = np.sum((y_pred - np.mean(y)) ** 2)
r2 = explained_variance / total_variance

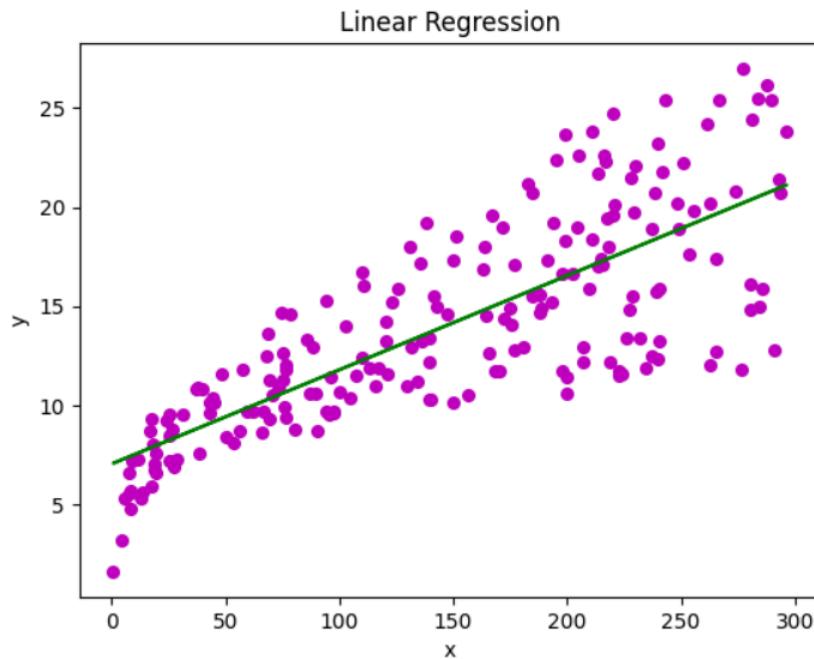
# Display results
print("Model Coefficients:", beta[1:].flatten())
print("Intercept:", beta[0][0])
print("Mean Squared Error:", mse)
print("R-squared Score:", r2)

# Plot actual vs predicted values
plt.scatter(y, y_pred, color='blue')
plt.plot(y, y, color='red', linestyle='--')
plt.xlabel("Actual Values")
plt.ylabel("Predicted Values")
plt.title("Actual vs Predicted Values")
plt.show()

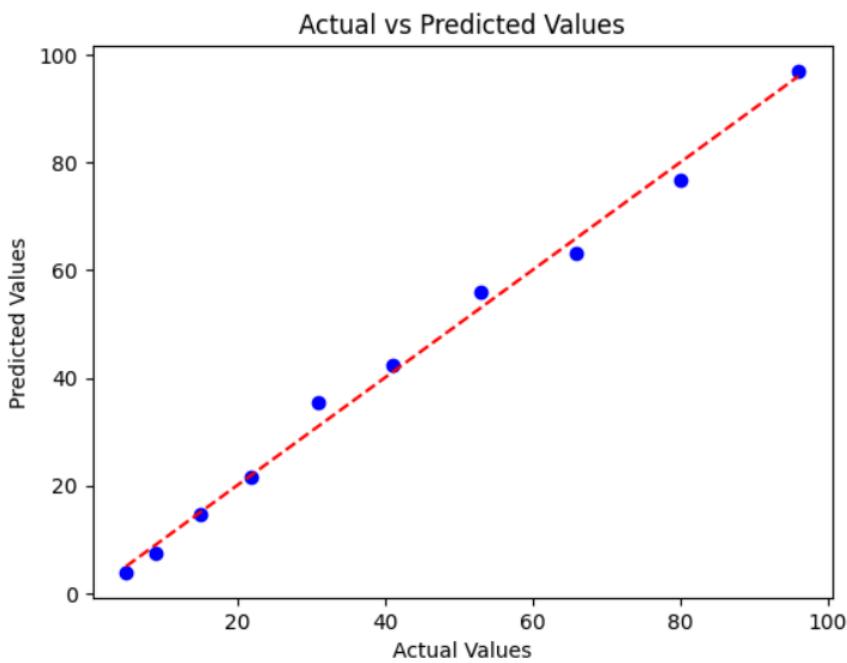
```

Output:

```
Enter the path to the CSV file: /content/tvmarketing.csv
Estimated coefficients:
b_0 = 7.0325935491276965
b_1 = 0.047536640433019736
```



```
Model Coefficients: [0.04026032 3.31385993 0.12078097]
Intercept: -3.1599509492879228
Mean Squared Error: 5.362912814290877
R-squared Score: 0.9935326149415827
```



Program 4

Build Logistic Regression Model for a given dataset

Screenshot:

```
Logistic Regression
def sigmoid(z):
    return 1 / (1 + np.exp(-z))

def compute(X, y, theta):
    m = len(y)
    h = sigmoid(X @ theta)
    cost = (-1/m) * np.sum(y * np.log(h) + (1-y) * np.log(1-h))
    return cost

def grad_descent(X, y, theta, alpha, iterations):
    m = len(y)
    cost_history = []
    for i in range(iterations):
        gradient = (1/m) * X.T @ (sigmoid(X @ theta) - y)
        theta -= alpha * gradient
    return theta, cost_history

def predict(X, theta):
    return (sigmoid(X @ theta) >= 0.5).astype(int)

accuracy = np.mean(y == predict(X, theta))
print("Accuracy: " + str(accuracy))

Output:
Accuracy: 1
```

Code:

```
#Logistic Regression

import numpy as np

import matplotlib.pyplot as plt

def sigmoid(z):

    return 1 / (1 + np.exp(-z))

def compute_cost(X, y, theta):

    m = len(y)

    h = sigmoid(X @ theta)

    cost = (-1/m) * np.sum(y * np.log(h) + (1 - y) * np.log(1 - h))

    return cost

def gradient_descent(X, y, theta, alpha, iterations):

    m = len(y)

    cost_history = []

    for _ in range(iterations):

        gradient = (1/m) * X.T @ (sigmoid(X @ theta) - y)

        theta -= alpha * gradient

        cost_history.append(compute_cost(X, y, theta))

    return theta, cost_history

def predict(X, theta):

    return (sigmoid(X @ theta) >= 0.5).astype(int)

# Generate synthetic binary classification data

np.random.seed(42)

X = np.random.rand(100, 1) # Feature values between 0 and 10
```

```

y = (X > 5).astype(int).ravel() #Label: 1 if X> 5, else 0

# Add intercept term

X_b = np.c_[np.ones((X.shape[0], 1)), X]

# Initialize parameters

theta = np.zeros(X_b.shape[1])

alpha = 0.1

iterations = 1000

# Train logistic regression using gradient descent

theta, cost_history = gradient_descent(X_b, y, theta, alpha, iterations)

# Make predictions

y_pred = predict(X_b, theta)

# Compute accuracy

accuracy = np.mean(y_pred == y)

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

# Plot the decision boundary

plt.scatter(X, y, color='blue', label='Actual Data')

plt.scatter(X, y_pred, color='red', marker='x', label='Predicted Labels')

plt.xlabel("Feature X")

plt.ylabel("Class (0 or 1)")

plt.legend()

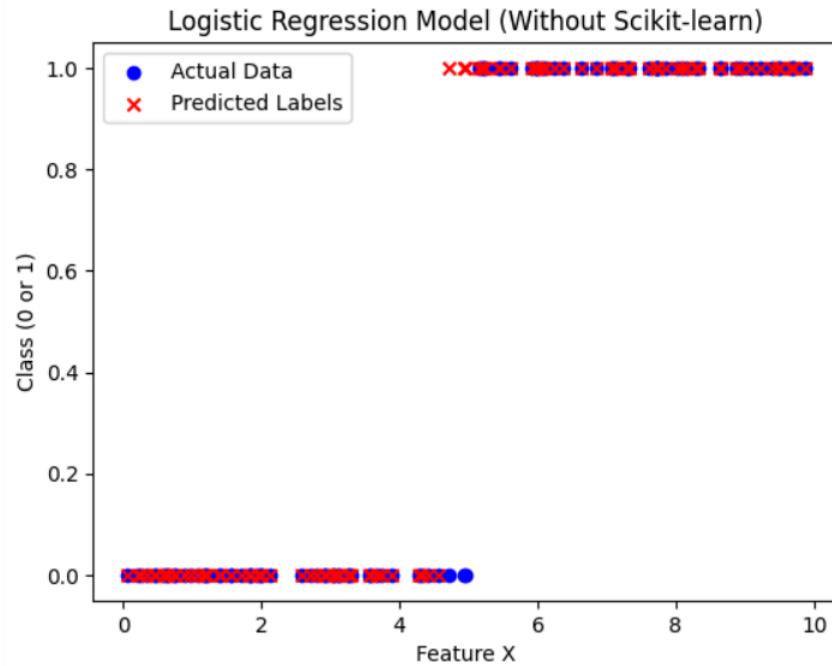
plt.title("Logistic Regression Model (Without Scikit-learn)")

plt.show()

```

Output:

Accuracy: 0.97



Program 5

Use an appropriate data set for building the decision tree (ID3) and apply this knowledge to classify a new sample

Screenshot:

The image shows handwritten Python code for the ID3 algorithm. The code is organized into several functions:

- Imports:** numpy, matplotlib.pyplot, pandas, math, copy.
- Dataset Loading:** dataset = pd.read_csv('content/weather.csv').
X = dataset.iloc[:, :].values.
X = X.astype('float')
- Attribute Selection:** attribute = ['outlook', 'Temp', 'Humidity', 'wind']
- Node Class Definition:** class Node(object):
def __init__(self):
 self.value = None
 self.decision = None
 self.child = None
- Entropy Calculation:** def findEntropy(data, rows):
 yes = 0
 no = 0
 ans = -1
 idx = len(data[0]) - 1
 entropy = 0

 for i in rows:
 if data[i][idx] == 'yes':
 yes += 1
 else:
 no += 1

 x = yes / (yes + no)
 y = no / (yes + no)
 if x != 0 and y != 0:
 entropy = -1 * (x * math.log2(x) + y * math.log2(y))
- Decision Tree Building:** if x == 1:
 ans = 1
if y == 1:
 ans = 0
return entropy, ans

```

def findMaxGain(data, rows, columns):
    maxGain = 0
    retIdx = -1
    entropy, ans = findEntropy(data, rows)
    if entropy == 0:
        return maxGain, retIdx, ans
    for j in columns:
        mydict = {}
        idx = j
        for i in rows:
            key = data[i][idx]
            if key not in mydict:
                mydict[key] = 1
            else:
                mydict[key] = mydict[key] + 1
        gain = entropy - mydict[key] * math.log2(key)
        if gain > maxGain:
            maxGain = gain
            retIdx = j
    return maxGain, retIdx, ans

```

```

def buildtree(data, rows, columns):
    margin, idx, ans = findMargin(Y, rows, columns)
    root = Node()
    root.childs = []
    if margin == 0:
        if ans == 1:
            root.value = 'Yes'
        else:
            root.value = 'No'
    return root

root.value = attribute[idx]
mydict = {}
for i in rows:
    key = data[i][idx]
    if key not in mydict:
        mydict[key] = 1
    else:
        mydict[key] += 1
newcolumns = copy.deepcopy(columns)
newcolumns.remove(idx)
for key in mydict:
    newrows = []
    for i in rows:
        if data[i][idx] == key:
            newrows.append(i)
    temp = buildTree(data, newrows, newcolumns)
    temp.decision = key
    root.childs.append(temp)
return root

def traverse(root, level=0):
    indent = " " * level
    print(f'{indent} Decision: {root.decision}, value: {root.value}')
    for i, child in enumerate(root.childs):
        if i == len(root.childs) - 1:
            traverse(child, level+1)
        else:
            print(" " * level + " ")

```

Code:

```
import numpy as np  
  
import matplotlib.pyplot as plt  
  
import pandas as pd  
  
import math  
  
import copy  
  
dataset = pd.read_csv('/content/weather.csv')  
  
X = dataset.iloc[:, :].values  
  
X  
  
attribute = ['Outlook', 'Temp', 'Humidity', 'Wi']
```

```

class Node(object):

    def __init__(self):
        self.value = None
        self.decision = None
        self.child = None

    def findEntropy(data, rows):
        yes=0
        no=0
        ans=-1
        idx=len(data[0])-1
        entropy=0
        for i in rows:
            if data[i][idx]=='Yes':
                yes=yes+1
            else:
                no=no+1
        x=yes/(yes+no)
        y=no/(yes+no)
        if x!=0 and y!=0:
            entropy= -1*(x*math.log2(x)+y*math.log2(y))
        if x==1:
            ans = 1
        if y==1:
            ans = 0

```

```

return entropy, ans

def findMaxGain(data, rows, columns):
    maxGain = 0
    retidx = -1
    entropy, ans = findEntropy(data, rows)
    if entropy == 0:
        """if ans == 1:
            print("Yes")
        else:
            print("No")"""
    return maxGain, retidx, ans

for j in columns:
    mydict = {}
    idx = j
    for i in rows:
        key = data[i][idx]
        if key not in mydict:
            mydict[key] = 1
        else:
            mydict[key] = mydict[key] + 1
    gain = entropy
    # print(mydict)
    for key in mydict:
        yes = 0

```

```

no = 0

for k in rows:

    if data[k][j] == key:

        if data[k][-1] == 'Yes':

            yes = yes + 1

        else:

            no = no + 1

    # print(yes, no)

    x = yes/(yes+no)

    y = no/(yes+no)

    # print(x, y)

    if x != 0 and y != 0:

        gain += (mydict[key] * (x*math.log2(x) + y*math.log2(y)))/14

    # print(gain)

    if gain > maxGain:

        # print("hello")

        maxGain = gain

        retidx = j

return maxGain, retidx, ans

def buildTree(data, rows, columns):

    maxGain, idx, ans = findMaxGain(X, rows, columns)

    root = Node()

    root.childs = []

    # print(maxGain)

```

```

if maxGain == 0:
    if ans == 1:
        root.value = 'Yes'
    else:
        root.value = 'No'
    return root

root.value = attribute[idx]
mydict = {}

for i in rows:
    key = data[i][idx]
    if key not in mydict:
        mydict[key] = 1
    else:
        mydict[key] += 1

newcolumns = copy.deepcopy(columns)
newcolumns.remove(idx)

for key in mydict:
    newrows = []
    for i in rows:
        if data[i][idx] == key:
            newrows.append(i)
    #print(newrows)
    temp = buildTree(data, newrows, newcolumns)

```

```

temp.decision = key

root.childs.append(temp)

return root

def traverse(root, level=0):

    # Create indentation based on the level

    indent = "    " * level # 4 spaces per level

    # Print the decision and value of the current node with indentation

    print(f"{indent} |--- Decision: {root.decision}, Value: {root.value}")

    # Traverse the children, if they exist

    for i, child in enumerate(root.childs):

        if i == len(root.childs) - 1: # Last child

            traverse(child, level + 1)

        else:

            traverse(child, level + 1)

def calculate():

    rows = [i for i in range(0, 14)]

    columns = [i for i in range(0, 4)]

    root = buildTree(X, rows, columns)

    root.decision = 'Start'

    traverse(root)

    calculate()

from graphviz import Source

dot_code = """

digraph G {

```

```

edge [dir=forward]

node [shape=box, style=bold]

A [label="OUTLOOK"]

B [label="HUMIDITY"]

C [label="WIND"]

D [label="NO", shape=plaintext]

E [label="YES", shape=plaintext]

F [label="YES", shape=plaintext]

G [label="NO", shape=plaintext]

A -> B [label="SUNNY"]

A -> E [label="OVERCAST"]

A -> C [label="RAIN"]

B -> D [label="HIGH"]

B -> F [label="NORMAL"]

C -> F [label="WEAK"]

C -> G [label="STRONG"]

}

"""

s = Source(dot_code, filename="decision_tree", format="png")

s.view()

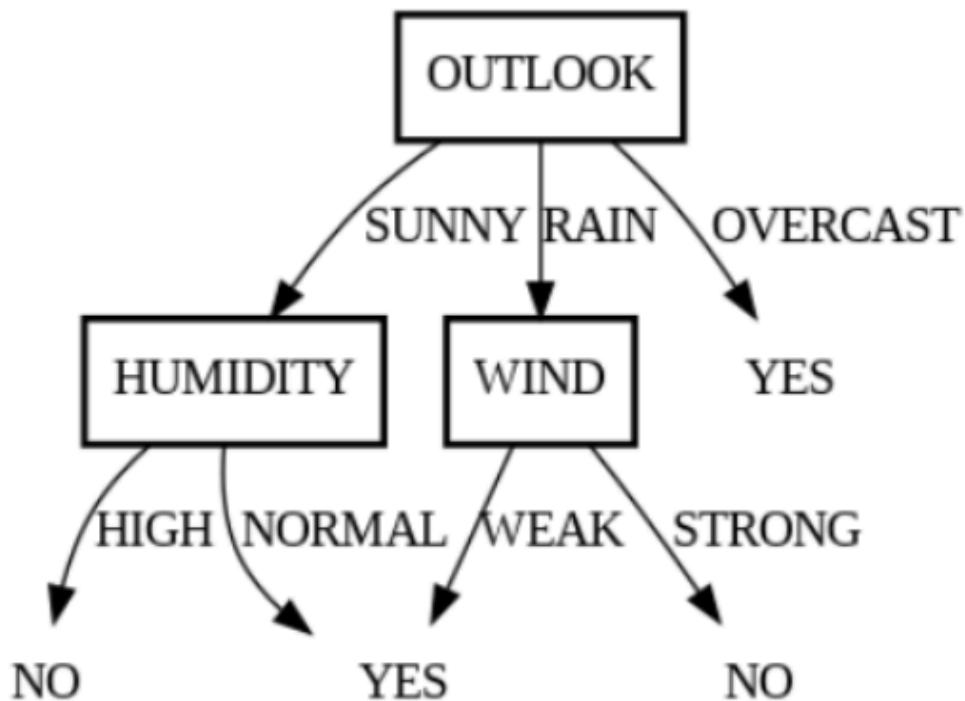
```

Output:

```
array([['Sunny', 'Hot', 'High', 'Weak', 'No'],
      ['Sunny', 'Hot', 'High', 'Strong', 'No'],
      ['Overcast', 'Hot', 'High', 'Weak', 'Yes'],
      ['Rain', 'Mild', 'High', 'Weak', 'Yes'],
      ['Rain', 'Cool', 'Normal', 'Weak', 'Yes'],
      ['Rain', 'Cool', 'Normal', 'Strong', 'No'],
      ['Overcast', 'Cool', 'Normal', 'Strong', 'Yes'],
      ['Sunny', 'Mild', 'High', 'Weak', 'No'],
      ['Sunny', 'Cool', 'Normal', 'Weak', 'Yes'],
      ['Rain', 'Mild', 'Normal', 'Weak', 'Yes'],
      ['Sunny', 'Mild', 'Normal', 'Strong', 'Yes'],
      ['Overcast', 'Mild', 'High', 'Strong', 'Yes'],
      ['Overcast', 'Hot', 'Normal', 'Weak', 'Yes'],
      ['Rain', 'Mild', 'High', 'Strong', 'No']], dtype=object)
```

```
|--- Decision: Start, Value: Outlook
|   |--- Decision: Sunny, Value: Humidity
|   |   |--- Decision: High, Value: No
|   |   |--- Decision: Normal, Value: Yes
|   |--- Decision: Overcast, Value: Yes
|   |--- Decision: Rain, Value: Wind
|   |   |--- Decision: Weak, Value: Yes
|   |   |--- Decision: Strong, Value: No
```

decision_tree.png X



Program 6

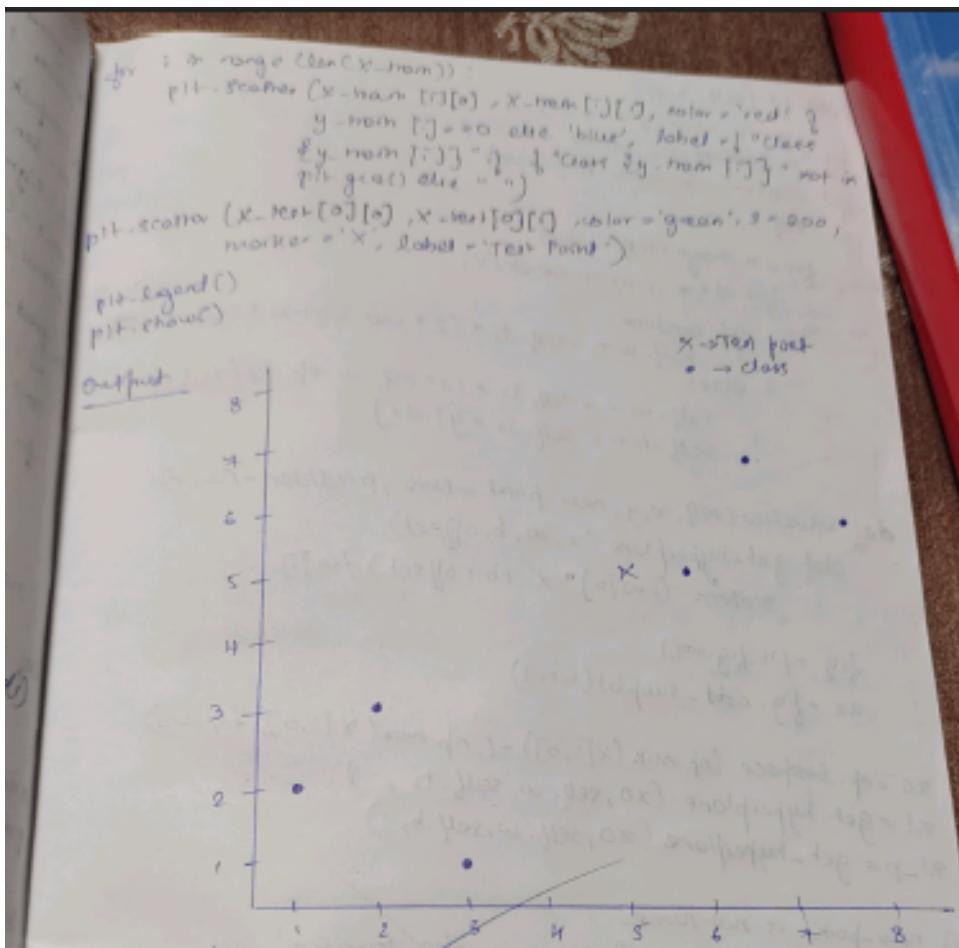
Build KNN Classification model for a given dataset

Screenshot:

The image shows handwritten code for a KNN classifier. The code is organized into several functions:

- KNN Algorithm:** Imports numpy and matplotlib.pyplot, defines a Counter class from collections, and implements a euclidean_distance function.
- Class KNN:** Contains methods for initializing (with k=3), fitting (X, y), predicting (X), and scoring (X, y). It uses np.argsort to find the k-nearest neighbors and np.mean to calculate the most common class.
- Data Preparation:** Shows the creation of X_train, y_train, X_test, and y_test arrays from lists of lists.
- Execution:** Shows the creation of a knn object, its fit method being called with X_train and y_train, and its predict method being called with X_test.
- Visualization:** Shows plt.figure(figsize=(8, 6)) which is likely intended to be part of the visualization code.

```
LAB-8  
KNN Algorithm:  
import numpy as np  
import matplotlib.pyplot as plt  
from collections import Counter  
def euclidean_distance(x1, x2):  
    return np.sqrt(np.sum((x1 - x2) ** 2))  
  
class KNN:  
    def __init__(self, k=3):  
        self.k = k  
    def fit(self, X, y):  
        self.X_train = np.array(X)  
        self.y_train = np.array(y)  
    def predict(self, X):  
        distances = [euclidean_distance(x, X_train) for x in X]  
        k_indices = np.argsort(distances)[:self.k]  
        k_nearest_labels = self.y_train[k_indices]  
        most_common = Counter(k_nearest_labels).most_common()  
        return most_common[0][0]  
    def score(self, X, y):  
        predictions = self.predict(X)  
        return np.mean(predictions == y)  
  
X_train = np.array([[1, 2], [2, 3], [3, 1], [6, 5], [7, 7], [8, 6]])  
y_train = np.array([0, 0, 0, 1, 1, 1])  
X_test = np.array([[5, 5]])  
  
knn = KNN(k=3)  
knn.fit(X_train, y_train)  
prediction = knn.predict(X_test)  
  
plt.figure(figsize=(8, 6))
```



Code:

```

import numpy as np

import matplotlib.pyplot as plt

from collections import Counter

```

Euclidean distance function

```

def euclidean_distance(x1, x2):
    return np.sqrt(np.sum((x1 - x2) ** 2))

```

KNN Class

class KNN:

```

    def __init__(self, k=3):

```

```

self.k = k

def fit(self, X, y):
    self.X_train = np.array(X)
    self.y_train = np.array(y)

def predict(self, X):
    return [self._predict(x) for x in X]

def _predict(self, x):
    distances = [euclidean_distance(x, x_train) for x_train in self.X_train]
    k_indices = np.argsort(distances)[:self.k]
    k_nearest_labels = [self.y_train[i] for i in k_indices]
    most_common = Counter(k_nearest_labels).most_common(1)
    return most_common[0][0]

def score(self, X, y):
    predictions = self.predict(X)
    return np.mean(predictions == y)

# Sample data

X_train = np.array([[1, 2], [2, 3], [3, 1], [6, 5], [7, 7], [8, 6]])
y_train = np.array([0, 0, 0, 1, 1, 1])
X_test = np.array([[5, 5]])

# Instantiate and fit KNN

knn = KNN(k=3)

knn.fit(X_train, y_train)

prediction = knn.predict(X_test)

```

```

# Plotting

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

# Training data

for i in range(len(X_train)):

    plt.scatter(X_train[i][0], X_train[i][1],
                color='red' if y_train[i] == 0 else 'blue',
                label=f"Class {y_train[i]}" if f"Class {y_train[i]}" not in
                plt.gca().get_legend_handles_labels()[1] else "")

# Test point

plt.scatter(X_test[0][0], X_test[0][1], color='green', s=200, marker='X', label='Test Point')

plt.title(f"KNN Classification (Predicted Class: {prediction[0]}"))

plt.xlabel("Feature 1")

plt.ylabel("Feature 2")

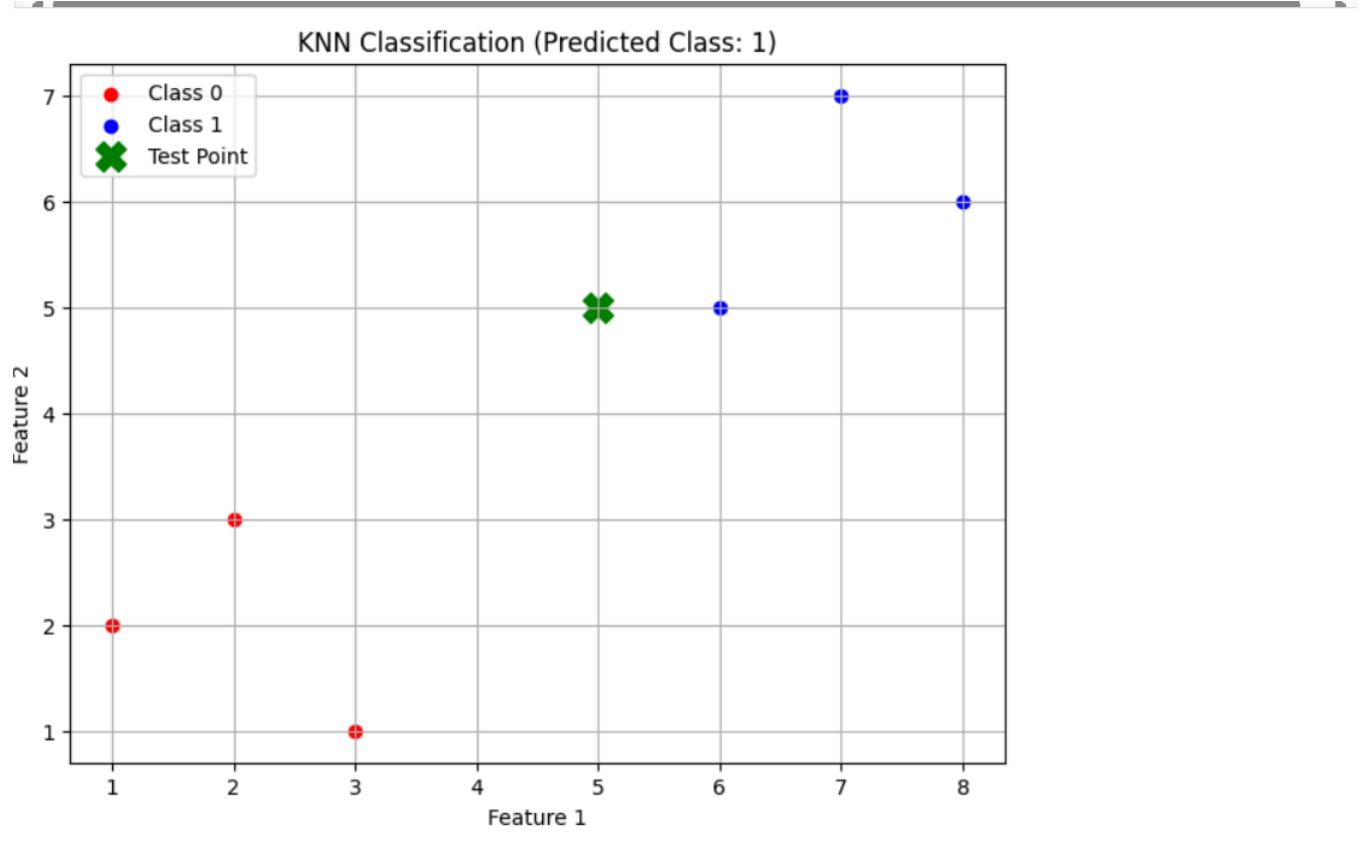
plt.legend()

plt.grid(True)

plt.show()

```

Output:



Program 7

Build Support vector machine model for a given dataset

Screenshot:

SVM algorithm

```

import numpy as np
import matplotlib.pyplot as plt

class SVM:
    def __init__(self, learning_rate=0.001, lambda_param=0.01):
        self.learning_rate = learning_rate
        self.lambda_param = lambda_param
        self.n_iters = n_iters
        self.w = None
        self.b = None
    
```

def fit(self, X, y):
 y = np.where(y <= 0, -1, 1)
 self.w = np.zeros(X.shape[1])
 self.b = 0
 for i in range(len(self.w)):
 for id_x, x_i in enumerate(X):
 if condition:
 self.w[i] += (2 * self.b + self.w * x_i) * y[id_x]
 else:
 self.w[i] += (2 * self.b + self.w * x_i) * -y[id_x]
 self.b += y[id_x]

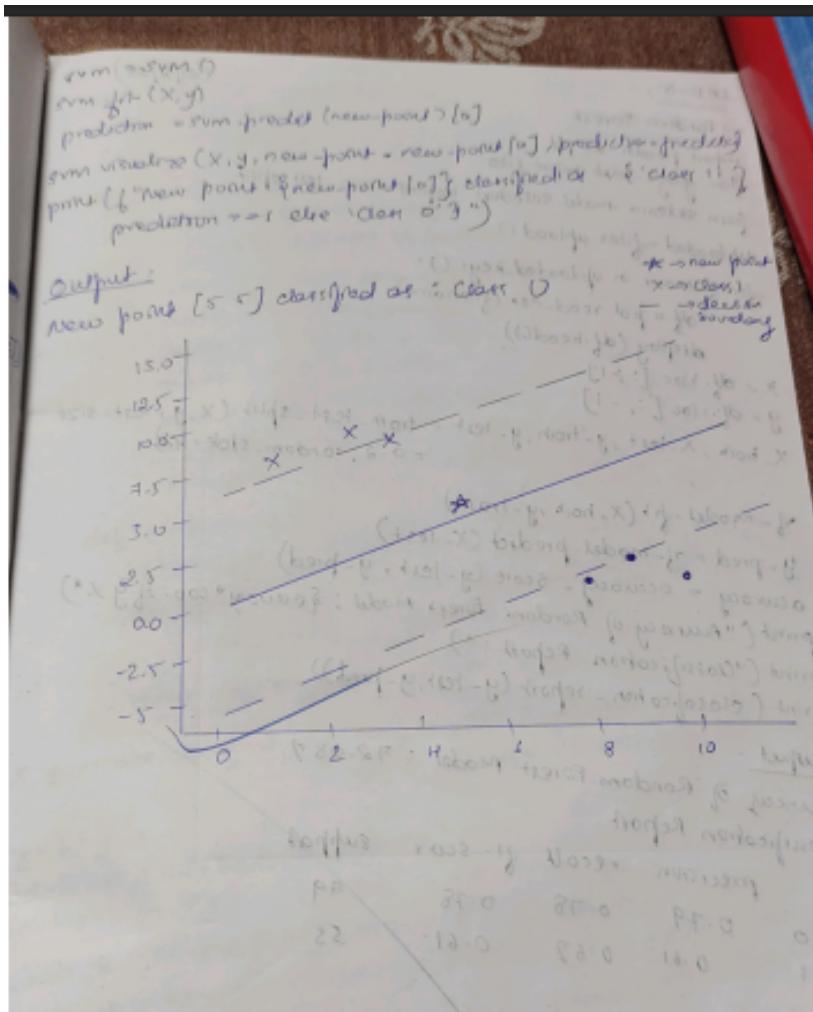
def visualize(self, x, y, new_point=None, prediction=None):
 def get_hyperplane(x, w, b, offset):
 return (-w[0] * x + b + offset) / w[1]

~~fig = plt.figure()~~
~~ax = fig.add_subplot(1, 1, 1)~~
~~x0 = np.linspace(np.min(x[:, 0]) - 1, np.max(x[:, 0]) + 1, 100)~~
~~ax = get_hyperplane(x0, self.w, self.b, -1)~~
~~ax = get_hyperplane(x0, self.w, self.b, 1)~~

1) new_point is not None:
 color = 'green' if prediction == 1 else 'orange'
 label = f'new point: class {y[0]} if prediction == {y[0]}'
 plt.scatter(new_point[0], new_point[1], label=label)
 plt.show()

example

B --name-- == "main" --
 X = np.array([-1, 7, 12, 8, 3, 8, 18, 0, 9, 2])
 y = np.array([0, 0, 0, 1, 1, 1])
 new_point = np.array([5, 5])



Code:

```
import numpy as np
```

```
import matplotlib.pyplot as plt
```

class SVM:

```
def __init__(self, learning_rate=0.001, lambda_param=0.01, n_iters=1000):  
    self.lr = learning_rate  
    self.lambda_param = lambda_param  
    self.n_iters = n_iters
```

```

self.w = None
self.b = None

def fit(self, X, y):
    y = np.where(y <= 0, -1, 1) # Convert labels to -1 and 1

    n_samples, n_features = X.shape
    self.w = np.zeros(n_features)
    self.b = 0

    for _ in range(self.n_iters):
        for idx, x_i in enumerate(X):
            condition = y[idx] * (np.dot(x_i, self.w) + self.b) >= 1

            if condition:
                self.w -= self.lr * (2 * self.lambda_param * self.w)
            else:
                self.w -= self.lr * (2 * self.lambda_param * self.w - np.dot(x_i, y[idx]))
                self.b += self.lr * y[idx]

def predict(self, X):
    approx = np.dot(X, self.w) + self.b
    return np.sign(approx)

def visualize(self, X, y, new_point=None, prediction=None):
    def get_hyperplane(x, w, b, offset):
        return (-w[0] * x + b + offset) / w[1]

    fig = plt.figure()
    ax = fig.add_subplot(1, 1, 1)

    # Plotting the data points
    for i in range(n_samples):
        if y[i] == 1:
            plt.scatter(X[i][0], X[i][1], color='red')
        else:
            plt.scatter(X[i][0], X[i][1], color='blue')

    # Plotting the decision boundary
    x_min, x_max = plt.xlim()
    y_min, y_max = plt.ylim()
    x_line = np.linspace(x_min, x_max)
    y_line = get_hyperplane(x_line, self.w, self.b, 0)
    plt.plot(x_line, y_line, 'k-')

    # Plotting the margin lines
    margin_w = self.w[1]
    margin_b = self.b
    margin_y_min = (margin_w * x_min) + margin_b
    margin_y_max = (margin_w * x_max) + margin_b
    plt.plot([x_min, x_max], [margin_y_min, margin_y_max], 'k--')

    # Plotting the new point (if provided)
    if new_point is not None:
        plt.scatter(new_point[0], new_point[1], color='green', s=100)

    # Plotting the predicted point (if provided)
    if prediction is not None:
        plt.scatter(prediction[0], prediction[1], color='purple', s=100)

```

```

# Plot existing data points

for i, sample in enumerate(X):
    if y[i] == 1:
        plt.scatter(sample[0], sample[1], marker='o', color='blue', label='Class +1' if i == 0 else '')
    else:
        plt.scatter(sample[0], sample[1], marker='x', color='red', label='Class -1' if i == 0 else '')

# Plot decision boundary

x0 = np.linspace(np.min(X[:, 0])-1, np.max(X[:, 0])+1, 100)

x1 = get_hyperplane(x0, self.w, self.b, 0)
x1_m = get_hyperplane(x0, self.w, self.b, -1)
x1_p = get_hyperplane(x0, self.w, self.b, 1)

ax.plot(x0, x1, 'k-', label='Decision Boundary')
ax.plot(x0, x1_m, 'k--', label='Margins')
ax.plot(x0, x1_p, 'k--')

# Plot the new point

if new_point is not None:
    color = 'green' if prediction == 1 else 'orange'
    label = f'New Point: Class {"1" if prediction == 1 else "0"}'

    plt.scatter(new_point[0], new_point[1], c=color, s=100, edgecolors='black', label=label,
               marker='*')

ax.legend()
plt.xlabel("Feature 1")
plt.ylabel("Feature 2")
plt.title("SVM with New Point Prediction")
plt.grid(True)

```

```

plt.show()

# Example

if __name__ == "__main__":
    # Training data

    X = np.array([
        [1, 7],
        [2, 8],
        [3, 8],
        [8, 1],
        [9, 2],
        [10, 2]
    ])

    y = np.array([0, 0, 0, 1, 1, 1]) # 0 -> -I, 1 -> +I

    # New point to classify

    new_point = np.array([[5, 5]])

    # Train and predict

    svm = SVM()
    svm.fit(X, y)
    prediction = svm.predict(new_point)[0]

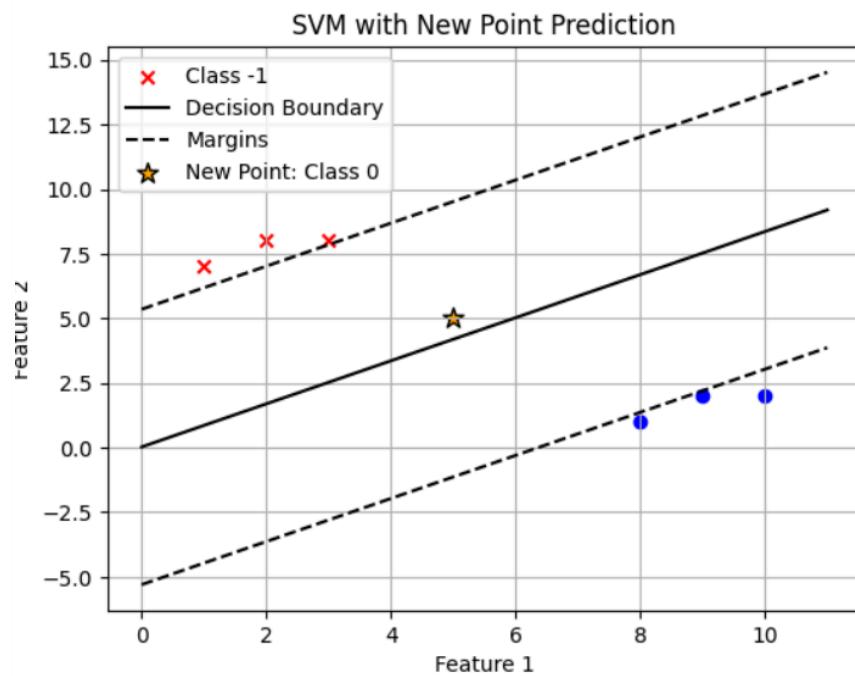
    # Visualize

    svm.visualize(X, y, new_point=new_point[0], prediction=prediction)

# Print prediction

print(f'New point {new_point[0]} classified as: {'Class 1' if prediction == 1 else 'Class 0'}")
```

Output:



New point [5 5] classified as: Class 0

Program 8

Implement Random forest ensemble method on a given dataset

Screenshot:

LAB - 5
(21/4/20)

```
→ Random Forest:  
import pandas as pd  
from google.colab import files  
from sklearn.model_selection import train_test_split  
uploaded = files.upload()  
for filename in uploaded.keys():  
    df = pd.read_csv(filename)  
    display(df.head())  
X = df.iloc[:, :-1]  
y = df.iloc[:, -1]  
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)  
rf_model = fit(X_train, y_train)  
y_pred = rf_model.predict(X_test)  
accuracy = accuracy_score(y_test, y_pred)  
print("Accuracy of Random Forest Model: {}%".format(accuracy * 100))  
print("Classification Report: ")  
print(classification_report(y_test, y_pred))  
  
Output:  
Accuracy of Random Forest Model: 72.08%  
classification Report:  
precision recall f1-score support  
0 0.79 0.78 0.78 99  
1 0.61 0.62 0.61 55
```

Code:

```
#RANDOM FOREST  
  
# STEP 1: Import required libraries  
import pandas as pd
```

```
from sklearn.model_selection import train_test_split  
from sklearn.ensemble import RandomForestClassifier  
from sklearn.metrics import accuracy_score, classification_report  
from google.colab import files
```

STEP 2: Upload your dataset

```
uploaded = files.upload()
```

STEP 3: Load the dataset (assuming it's a CSV)

```
for filename in uploaded.keys():  
    df = pd.read_csv(filename)  
    print(f'Data loaded from: {filename}')  
    display(df.head()) # Display first 5 rows of data
```

STEP 4: Preprocessing

Assume the last column is the target variable (label)

```
X = df.iloc[:, :-1] # Features (all rows, all columns except last)  
y = df.iloc[:, -1] # Target (last column)
```

STEP 5: Split the data into training and testing sets

```
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
```

STEP 6: Initialize and train the Random Forest model

```
rf_model = RandomForestClassifier(n_estimators=100, random_state=42) # 100 trees in the forest
```

```
rf_model.fit(X_train, y_train)
```

STEP 7: Make predictions on the test set

```
y_pred = rf_model.predict(X_test)
```

STEP 8: Evaluate the model

```
accuracy = accuracy_score(y_test, y_pred)
```

```
print(f'Accuracy of Random Forest Model: {accuracy * 100:.2f}%')
```

STEP 9: Print classification report

```
print("Classification Report:")
```

```
print(classification_report(y_test, y_pred))
```

Output:

Upload widget is only available when the cell has been executed in the current browser session. Please rerun this cell to enable.

Saving diabetes.csv to diabetes.csv

Data loaded from: diabetes.csv

Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	DiabetesPedigreeFunction	Age	Outcome
0	6	148	72	35	0	33.6	0.627	50
1	1	85	66	29	0	26.6	0.351	31
2	8	183	64	0	0	23.3	0.672	32
3	1	89	66	23	94	28.1	0.167	21
4	0	137	40	35	168	43.1	2.288	33

Accuracy of Random Forest Model: 72.08%

Classification Report:

	precision	recall	f1-score	support
0	0.79	0.78	0.78	99
1	0.61	0.62	0.61	55
accuracy			0.72	154
macro avg	0.70	0.70	0.70	154
weighted avg	0.72	0.72	0.72	154

Program 9

Implement Boosting ensemble method on a given dataset

Screenshot:

→ Boosting :

```
import numpy as np
import seaborn as sns
from sklearn.tree import DecisionTreeClassifier
sns.set(style = "whitegrid")
```

```
class AdaBoost:
    def __init__(self, n_estimators = 50):
        self.alpha = []
        self.models = []
        self.errors = []
    def fit(self, X, y):
        n_samples = X.shape[0]
        w = np.ones(n_samples) / n_samples
        for estimator in range(self.n_estimators):
            model = DecisionTreeClassifier(max_depth=1)
            model.fit(X, y, sample_weight=w)
            y_pred = model.predict(X)
            alpha = 0.5 * np.log((1 - err) / err)
            w = w * np.exp(-alpha * (y - y_pred))
            w = w / np.sum(w)
    def predict(self, X):
        final_pred = np.zeros(X.shape[0])
        for model, alpha in zip(self.models, self.alpha):
            final_pred += alpha * model.predict(X)
        return np.sign(final_pred)
```

X, y = make_classification(n_samples = 500, n_features = 2, n_info = 2, n_redundant = 0, n_classes = 2, random_state = 42)

y = 2 * y - 1

adaBoost = AdaBoost(n_estimators = 50)

adaBoost.fit(X, y)

x_min, x_max = X[:, 0].min() - 1, X[:, 0].max() + 1, c = "#000000"
y_min, y_max = X[:, 1].min() - 1, X[:, 1].max() + 1, c = "#000000"

z = adaBoost.predict(np.c_[x.ravel(), y.ravel()])

z = z.reshape(X.shape)

plt.figure(figsize=(10, 6))
plt.show()

(1) zoomed in around z
(2) zoomed in

Scatter plot of X vs y
("X" = sklearn.datasets.iris)

Code:

```
#BOOSTING

import numpy as np

import matplotlib.pyplot as plt

import seaborn as sns

from sklearn.tree import DecisionTreeClassifier

from sklearn.datasets import make_classification

from sklearn.metrics import accuracy_score

from sklearn.decomposition import PCA


# Set up plot style

sns.set(style="whitegrid")

class AdaBoost:

    def __init__(self, n_estimators=50):

        self.n_estimators = n_estimators

        self.alphas = [] # Weights of each weak classifier

        self.models = [] # Weak classifiers (e.g., decision stumps)

        self.errors = [] # List to store error for each estimator

    def fit(self, X, y):

        # Initialize weights for each data point

        n_samples, n_features = X.shape

        w = np.ones(n_samples) / n_samples # Equal weights initially

        for estimator in range(self.n_estimators):

            # Train weak classifier (decision stump)
```

```

model = DecisionTreeClassifier(max_depth=1) # Decision stump

model.fit(X, y, sample_weight=w)

y_pred = model.predict(X)

# Calculate error rate

err = np.sum(w * (y_pred != y)) / np.sum(w)

self.errors.append(err)

# Compute alpha (weight for the classifier)

alpha = 0.5 * np.log((1 - err) / err) if err < 1 else 0

self.alphas.append(alpha)

self.models.append(model)

# Update weights for misclassified samples

w = w * np.exp(-alpha * y * y_pred) # Update weights based on classifier performance

w = w / np.sum(w) # Normalize the weights


def predict(self, X):

    # Initialize the final prediction

    final_pred = np.zeros(X.shape[0])

    for model, alpha in zip(self.models, self.alphas):

        final_pred += alpha * model.predict(X)

    # Return the sign of the final prediction

    return np.sign(final_pred)

def score(self, X, y):

    # Return accuracy of the model

    return accuracy_score(y, self.predict(X))

```

```

# Generate a synthetic binary classification dataset with 2 informative features

X, y = make_classification(n_samples=500, n_features=2, n_informative=2, n_redundant=0,
n_classes=2, random_state=42)

#Convert labels to -1 and 1 for AdaBoost

y = 2 * y - 1

# Create and train AdaBoost model

adaboost = AdaBoost(n_estimators=50)

adaboost.fit(X, y)

# Evaluate the model

accuracy = adaboost.score(X, y)

print(f"Model accuracy: {accuracy:.4f}")

# Plot error over iterations

plt.figure(figsize=(10, 6))

plt.plot(range(1, adaboost.n_estimators + 1), adaboost.errors, marker='o', linestyle='-', color='b')

plt.title('Error vs. Number of Estimators')

plt.xlabel('Number of Estimators')

plt.ylabel('Error')

plt.grid(True)

plt.show()

# Plot decision boundary for final model

x_min, x_max = X[:, 0].min() - 1, X[:, 0].max() + 1

y_min, y_max = X[:, 1].min() - 1, X[:, 1].max() + 1

xx, yy = np.meshgrid(np.arange(x_min, x_max, 0.1),
                     np.arange(y_min, y_max, 0.1))

```

```

Z = adaboost.predict(np.c_[xx.ravel(), yy.ravel()])

Z = Z.reshape(xx.shape)

plt.figure(figsize=(10, 6))

plt.contourf(xx, yy, Z, alpha=0.75, cmap='coolwarm')

plt.scatter(X[:, 0], X[:, 1], c=y, edgecolors='k', marker='o', s=50, cmap='coolwarm')

plt.title('AdaBoost Decision Boundary')

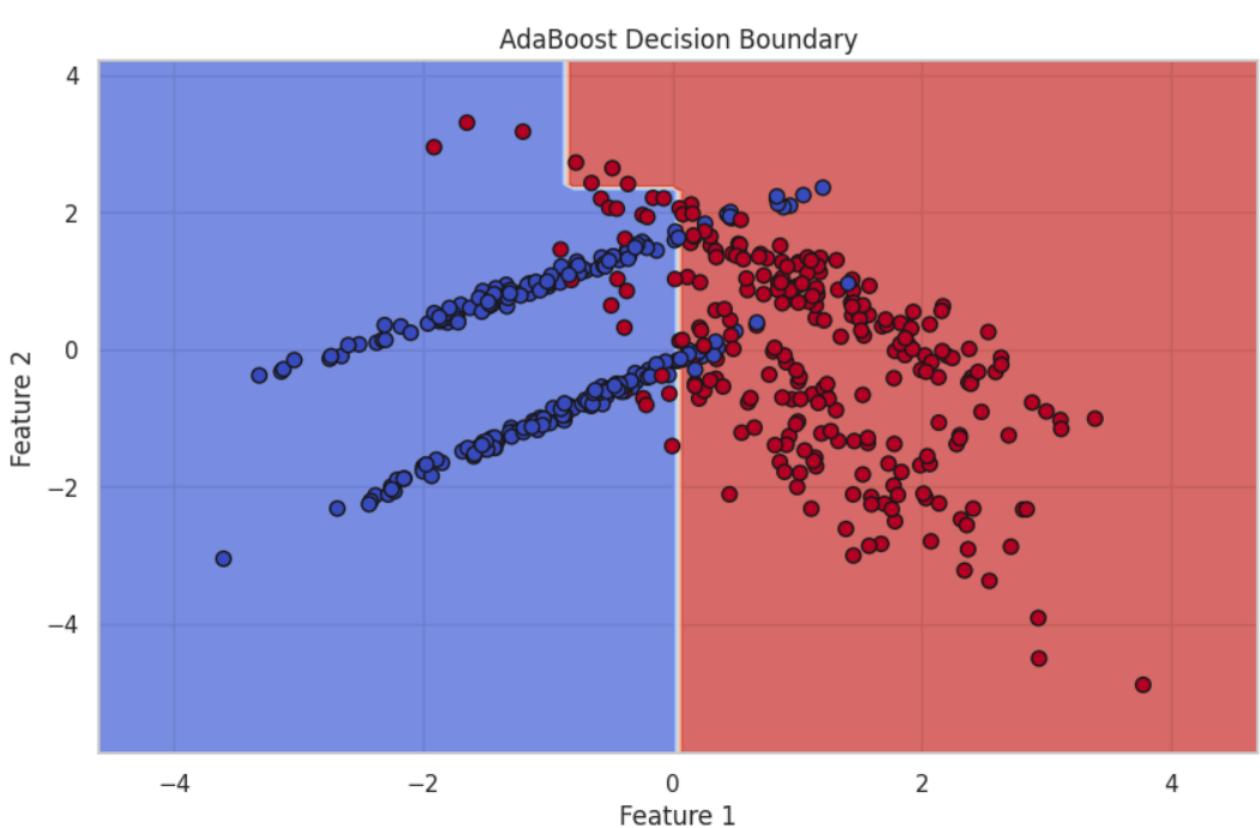
plt.xlabel('Feature 1')

plt.ylabel('Feature 2')

plt.show()

```

Output:



Program 10

Build k-Means algorithm to cluster a set of data stored in a .CSV file

Screenshot:

```

→ K-Means Clustering
class K_Means:
    def __init__(self, k=3, tolerance=0.001, max_iter=500):
        self.k = k
        self.tolerance = tolerance
        self.max_iter = max_iter
        self.centroids = None
        self.classifications = None
        self.colors = None
    def predict(self, data):
        self.classifications = np.zeros(len(data))
        self.colors = np.zeros(len(data))
        for i in range(self.max_iter):
            self.classifications = self.classify(data)
            self.colors = self.get_color(self.classifications)
            if self.converged():
                break
            self.centroids = self.get_centroids(data)
    def fit(self, data):
        for i in range(self.k):
            self.classifications[i] = data[i]
            self.colors[i] = np.random.randint(0, len(self.colors))
        for i in range(self.max_iter):
            self.classifications = self.classify(data)
            self.colors = self.get_color(self.classifications)
            if self.converged():
                break
            self.centroids = self.get_centroids(data)
    def main():
        K = 3
        center_1 = np.array([1, 1])
        center_2 = np.array([5, 5])
        center_3 = np.array([8, 1])
        cluster_1 = np.random.rand(100, 2) + center_1
        cluster_2 = np.random.rand(100, 2) + center_2
        cluster_3 = np.random.rand(100, 2) + center_3
        k_mean = K_Means(K)
        k_mean.fit(data)
        for centroid in k_mean.centroids:
            plt.scatter(s=130, marker="x")
        for cluster_index in k_mean.classifications:
            color = colors[cluster_index]
            for feature in k_mean.features[cluster_index]:
                plt.scatter(feature[0], feature[1], color=color, s=90)
        plt.show()

```

```

for cluster_index in k_mean.classifications:
    color = colors[cluster_index]
    for feature in k_mean.features[cluster_index]:
        plt.scatter(feature[0], feature[1], color=color, s=90)
    if None == "main__":
        main()
        analysis(PCA)

```

Code:

```
#K-MEANS

class K_Means:

    def __init__(self, k=2, tolerance = 0.001, max_iter = 500):
        self.k = k
        self.max_iterations = max_iter
        self.tolerance = tolerance

    def euclidean_distance(self, point1, point2):
        #return math.sqrt((point1[0]-point2[0])**2 + (point1[1]-point2[1])**2 +
        #                (point1[2]-point2[2])**2) #sqrt((x1-x2)^2 + (y1-y2)^2)
        return np.linalg.norm(point1-point2, axis=0)

    def predict(self,data):
        distances = [np.linalg.norm(data-self.centroids[centroid]) for centroid in self.centroids]
        classification = distances.index(min(distances))
        return classification

    def fit(self, data):
        self.centroids = {}
        for i in range(self.k):
            self.centroids[i] = data[i]

        for i in range(self.max_iterations):
            self.classes = {}
            for j in range(self.k):
                self.classes[j] = []
            for point in data:
                distances = []
                for centroid in self.centroids:
                    distances.append(self.euclidean_distance(point, self.centroids[centroid]))
                classification = distances.index(min(distances))
                self.classes[classification].append(point)
```

```

for index in self.centroids:

    distances.append(self.euclidean_distance(point,self.centroids[index]))

    cluster_index = distances.index(min(distances))

    self.classes[cluster_index].append(point)

previous = dict(self.centroids)

for cluster_index in self.classes:

    self.centroids[cluster_index] = np.average(self.classes[cluster_index], axis = 0)

isOptimal = True

for centroid in self.centroids:

    original_centroid = previous[centroid]

    curr = self.centroids[centroid]

    if np.sum((curr - original_centroid)/original_centroid * 100.0) > self.tolerance:

        isOptimal = False

    if isOptimal:

        break

def main():

    K=3

    center_1 = np.array([1,1])

    center_2 = np.array([5,5])

    center_3 = np.array([8,1])

    # Generate random data and center it to the three centers

    cluster_1 = np.random.randn(100, 2) + center_1

    cluster_2 = np.random.randn(100,2) + center_2

    cluster_3 = np.random.randn(100,2) + center_3

```

```

data = np.concatenate((cluster_1, cluster_2, cluster_3), axis = 0)

k_means = K_Means(K)

k_means.fit(data)

# Plotting starts here

colors = 10*["r", "g", "c", "b", "k"]

for centroid in k_means.centroids:

    plt.scatter(k_means.centroids[centroid][0], k_means.centroids[centroid][1], s = 130, marker = "x")

    for cluster_index in k_means.classes:

        color = colors[cluster_index]

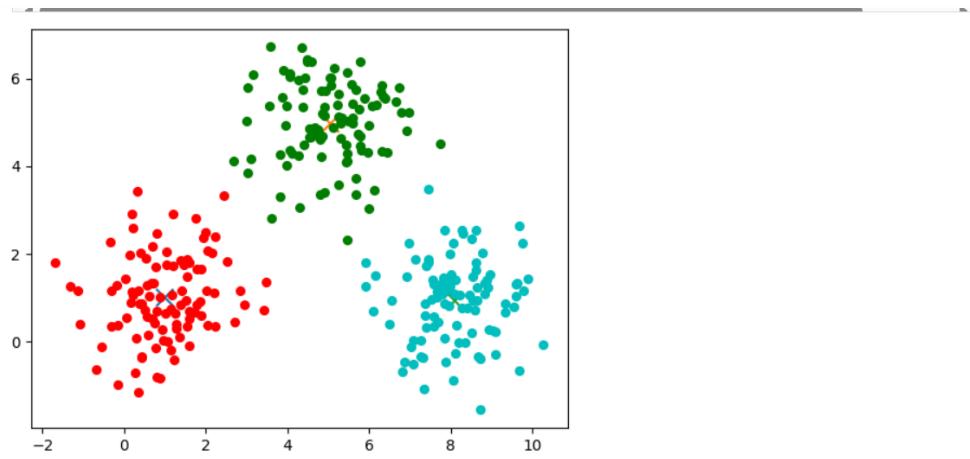
        for features in k_means.classes[cluster_index]:

            plt.scatter(features[0], features[1], color = color,s = 30)

if __name__ == "__main__":
    main()

```

Output:



Program 11

Implement Dimensionality reduction using Principal Component Analysis (PCA) method

Screenshot:

```
# main()
# Principle component analysis (PCA)
import pandas as pd
from sklearn.decomposition import PCA
uploaded = files.upload()
for filenam in uploaded:
    df = pd.read_csv(filenam)
    display(df.head())
numeric_df = df.select_dtypes(include = [np.number])
selected_features = numeric_df.columns
X = numeric_df[selected_features].dropna()
X_scaled = StandardScaler().fit_transform(X)
pca = PCA(n_components = 2)
principal_components = pca.fit_transform(X_scaled)
principal_components = pd.DataFrame(principal_components, columns = [f'PC{i}' for i in range(2)])
plt.figure(figsize=(6,6))
plt.gcf().show()
print("Explained variance Ratio : ", pca.explained_variance_ratio_)

Output:
Explained variance Ratio : [0.5216, 0.2863]
```

C. S. R. M.

Code:

```
#PCA
```

STEP 1: Import packages

```
import pandas as pd
```

```
import numpy as np
```

```
from sklearn.decomposition import PCA
```

```
from sklearn.preprocessing import StandardScaler
```

```
import matplotlib.pyplot as plt
```

```
from google.colab import files
```

STEP 2: Upload the CSV file

```
uploaded = files.upload()
```

STEP 3: Load the dataset

```
for filename in uploaded.keys():
```

```
    df = pd.read_csv(filename)
```

```
    print(f" Uploaded: {filename}")
```

```
    display(df.head())
```

STEP 4: Select numerical columns

```
numeric_df = df.select_dtypes(include=[np.number])
```

```
print("Numerical features found:", list(numeric_df.columns))
```

OPTIONAL: Manually select columns if needed

```

# selected_features = ['feature1', 'feature2', ...]

selected_features = numeric_df.columns # use all numeric features for now


# STEP 5: Standardize data

X = numeric_df[selected_features].dropna()

X_scaled = StandardScaler().fit_transform(X)


# STEP 6: Apply PCA

pca = PCA(n_components=2)

principal_components = pca.fit_transform(X_scaled)


# STEP 7: Create DataFrame for components

pca_df = pd.DataFrame(data=principal_components, columns=['PC1', 'PC2'])


# STEP 8: Visualize the first two principal components

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

plt.scatter(pca_df['PC1'], pca_df['PC2'], alpha=0.7)

plt.xlabel('Principal Component 1')

plt.ylabel('Principal Component 2')

plt.title('2D PCA Visualization')

plt.grid(True)

plt.show()


# STEP 9: Explained variance ratio

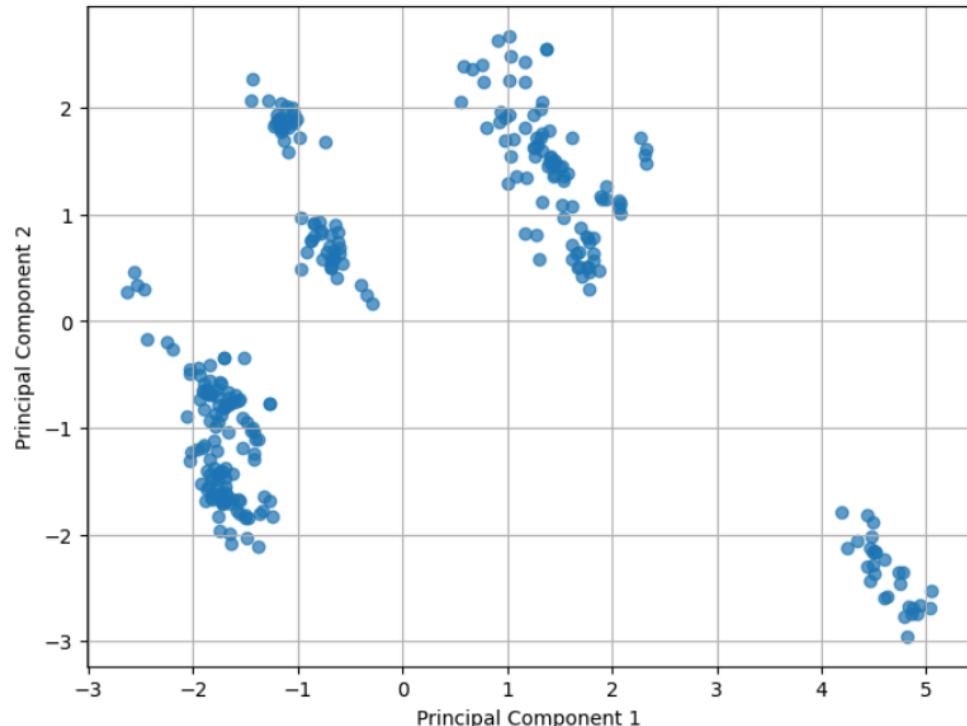
print("Explained Variance Ratio:", pca.explained_variance_ratio_)

```

Output:

```
NUM Numerical features found: ['id', 'mois', 'prot', 'fat', 'ash', 'sodium', 'carb', 'cal']
```

2D PCA Visualization



```
EXPL Explained Variance Ratio: [0.52163044 0.28631263]
```