

VISVESVARAYA TECHNOLOGICAL UNIVERSITY

“JnanaSangama”, Belgaum- 590014, Karnataka.



LAB REPORT on **Machine Learning (23CS6PCMAL)**

Submitted by

Likhith M (1BM22CS135)

in partial fulfillment for the award of the degree of

BACHELOR OF ENGINEERING

in

COMPUTER SCIENCE AND 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 **Likhith M (1BM22CS135)**, who is bonafide student of **B.M.S. College of Engineering**. It is in partial fulfilment for the award of **Bachelor of Engineering in Computer Science and Engineering** of the Visvesvaraya Technological University, Belgaum. The Laboratory report has been approved as it satisfies the academic requirements in respect of an Machine Learning (23CS6PCMAL) work prescribed for the said degree.

Sheetal V A
Assistant Professor
Department of CSE, BMSCE

Dr. Kavitha Sooda
Professor & HOD
Department of CSE, BMSCE

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Github Link: <https://github.com/01-BLUELOTUS/ML-1BM22CS135.git>

LABORATORY PROGRAM – 1

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

OBSERVATION BOOK

2.3.25 Date _____
Page _____

LABORATORY PROGRAM - 1
Write a python program to Import and export data using Pandas library functions.

```
import pandas as pd  
df = pd.read_csv('housing.csv')  
print("Dataframe Information :")  
df.info()  
  
print("In Statistical Summary :")  
print(df.describe())  
  
print("In Unique Value Counts for 'Ocean Proximity':")  
print(df['Ocean Proximity'].value_counts())  
  
print("In Columns with missing values :")  
missing_counts = df.isnull().sum()  
print(missing_counts[missing_counts > 0])
```

- After applying data processing techniques to Diabetes and Adult Income data sets:
- Which columns in the dataset had missing values? How did you handle them?
→ ~~print(diabetes_df.isnull().sum())~~
The code checks for missing values (NaN) in each column of the diabetes dataset and prints the count of missing values for each column. Numerical columns are filled with the mean of the column, and categorical columns with the mode.
2. Which categorical columns did you identify in the dataset? How do you encode them?
→ cat_cols_adult = adult_income_df.select_dtypes(include=[object]).columns.to_list()
This part identifies categorical columns in the adult income dataset. One-hot encoding is then applied to these columns using pd.get_dummies.
3. What is the difference between Min-Max scaling and Standardization? When would you use one over the other?
→ Min-Max scaling scales the data to a specific range (usually 0 to 1). It's sensitive to outliers.
• Standardization (Z-score normalization) transforms data to have a mean of 0 and a standard deviation of 1. It's less sensitive to outliers.
• Min-Max scaling is preferred when the algorithm is sensitive to the "magnitude" of the features (eg. KNN, K-Means). Use it when you know the data's distribution doesn't have significant outliers.
• Standardization is generally preferred if the data has a Gaussian-like distribution or when the algorithm is not sensitive to the features scales (eg. Linear regression, logistic regression). Use it when you're concerned about outliers influencing the scaling. Standardization is often more robust.

CODE WITH OUTPUT

Diabetes Dataset

```
df=pd.read_csv('/content/Dataset of Diabetes .csv')
df.head()
```

	ID	No_Pation	Gender	AGE	Urea	Cr	HbA1c	Chol	TG	HDL	LDL	VLDL	BMI	CLASS
0	502	17975	F	50	4.7	46	4.9	4.2	0.9	2.4	1.4	0.5	24.0	N
1	735	34221	M	26	4.5	62	4.9	3.7	1.4	1.1	2.1	0.6	23.0	N
2	420	47975	F	50	4.7	46	4.9	4.2	0.9	2.4	1.4	0.5	24.0	N
3	680	87656	F	50	4.7	46	4.9	4.2	0.9	2.4	1.4	0.5	24.0	N
4	504	34223	M	33	7.1	46	4.9	4.9	1.0	0.8	2.0	0.4	21.0	N

```
df.shape
```

```
(1000, 14)
```

```
print(df.info())
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 1000 entries, 0 to 999
Data columns (total 14 columns):
 #   Column      Non-Null Count  Dtype  
--- 
 0   ID          1000 non-null    int64  
 1   No_Pation   1000 non-null    int64  
 2   Gender      1000 non-null    object  
 3   AGE         1000 non-null    int64  
 4   Urea        1000 non-null    float64 
 5   Cr          1000 non-null    int64  
 6   HbA1c       1000 non-null    float64 
 7   Chol        1000 non-null    float64 
 8   TG          1000 non-null    float64 
 9   HDL         1000 non-null    float64 
 10  LDL         1000 non-null    float64 
 11  VLDL        1000 non-null    float64 
 12  BMI         1000 non-null    float64 
 13  CLASS        1000 non-null    object  
dtypes: float64(8), int64(4), object(2)
memory usage: 109.5+ KB
```

M None

```

# Summary statistics
print(df.describe())

      ID  No_Pation      AGE     Urea      Cr \
count 1000.000000 1.000000e+03 1000.000000 1000.000000 1000.000000 \
mean 340.500000 2.70514e+05 53.528000 5.124743 68.943000
std 240.397673 3.380758e+06 8.799241 2.935165 59.984747
min 1.000000 1.230000e+02 20.000000 0.500000 6.000000
25% 125.750000 2.406375e+04 51.000000 3.700000 48.000000
50% 300.500000 3.439550e+04 55.000000 4.600000 60.000000
75% 550.250000 4.538425e+04 59.000000 5.700000 73.000000
max 800.000000 7.543566e+07 79.000000 38.900000 800.000000

      HbA1c      Chol      TG      HDL      LDL \
count 1000.000000 1000.000000 1000.000000 1000.000000 1000.000000 \
mean 8.281160 4.862820 2.349610 1.204750 2.609790
std 2.534003 1.301738 1.401176 0.660414 1.115102
min 0.900000 0.000000 0.300000 0.200000 0.300000
25% 6.500000 4.000000 1.500000 0.900000 1.800000
50% 8.000000 4.800000 2.000000 1.100000 2.500000
75% 10.200000 5.600000 2.900000 1.300000 3.300000
max 16.000000 10.300000 13.800000 9.900000 9.900000

      VLDL      BMI
count 1000.000000 1000.000000
mean 1.854700 29.578020
std 3.663599 4.962388
min 0.100000 19.000000
25% 0.700000 26.000000
50% 0.900000 30.000000
75% 1.500000 33.000000
max 35.000000 47.750000

missing_values=df.isnull().sum()
print(missing_values[missing_values > 0])

Series([], dtype: int64)

categorical_cols = df.select_dtypes(include=['object']).columns
print("Categorical columns identified:", categorical_cols)
if len(categorical_cols) > 0:
    df = pd.get_dummies(df, columns=categorical_cols, drop_first=True)
    print("\nDataFrame after one-hot encoding:")
    print(df.head())
else:
    print("\nNo categorical columns found in the dataset.")

Categorical columns identified: Index(['Gender', 'CLASS'], dtype='object')

DataFrame after one-hot encoding:
      ID  No_Pation  AGE  Urea  Cr  HbA1c  Chol  TG  HDL  LDL  VLDL  BMI \
0   502       17975  50   4.7  46   4.9   4.2  0.9  2.4  1.4   0.5  24.0
1   735       34221  26   4.5  62   4.9   3.7  1.4  1.1  2.1   0.6  23.0
2   420       47975  50   4.7  46   4.9   4.2  0.9  2.4  1.4   0.5  24.0
3   680       87656  50   4.7  46   4.9   4.2  0.9  2.4  1.4   0.5  24.0
4   504       34223  33   7.1  46   4.9   4.9  1.0  0.8  2.0   0.4  21.0

      Gender_M  Gender_f  CLASS_N  CLASS_P  CLASS_Y  CLASS_Z \
0     False    False    False    False    False    False
1     True     False    False    False    False    False
2    False    False    False    False    False    False
3    False    False    False    False    False    False
4     True    False    False    False    False    False

```

```
from sklearn.preprocessing import MinMaxScaler, StandardScaler
import pandas as pd

numerical_cols = df.select_dtypes(include=['number']).columns

scaler = MinMaxScaler()
df_minmax = df.copy() # Create a copy to avoid modifying the original
df_minmax[numerical_cols] = scaler.fit_transform(df[numerical_cols])

scaler = StandardScaler()
df_standard = df.copy()
df_standard[numerical_cols] = scaler.fit_transform(df[numerical_cols])
print("\nDataFrame after Min-Max Scaling:")
print(df_minmax.head())
print("\nDataFrame after Standardization:")
print(df_standard.head())

DataFrame after Min-Max Scaling:
   ID No_Patien AGE    Urea     Cr HbA1c   Chol \
0  0.627034  0.000237 0.508475  0.109375  0.050378 0.264901  0.407767
1  0.918648  0.000452 0.101695  0.104167  0.070529 0.264901  0.359223
2  0.524406  0.000634 0.508475  0.109375  0.050378 0.264901  0.407767
3  0.849812  0.001160 0.508475  0.109375  0.050378 0.264901  0.407767
4  0.629537  0.000452 0.220339  0.171875  0.050378 0.264901  0.475728

      TG      HDL      LDL      VLDL      BMI Gender_M Gender_f \
0  0.044444  0.226804  0.114583  0.011461  0.173913 False   False
1  0.081481  0.092784  0.187500  0.014327  0.139130 True   False
2  0.044444  0.226804  0.114583  0.011461  0.173913 False   False
3  0.044444  0.226804  0.114583  0.011461  0.173913 False   False
4  0.051852  0.061856  0.177083  0.008596  0.069565 True   False

  CLASS_N CLASS_P CLASS_Y CLASS_Y
0  False   False   False   False
1  False   False   False   False
2  False   False   False   False
3  False   False   False   False
4  False   False   False   False

DataFrame after Standardization:
   ID No_Patien AGE    Urea     Cr HbA1c   Chol \
0  0.672140 -0.074747 -0.401144 -0.144781 -0.382672 -1.334983 -0.509436
1  1.641852 -0.069940 -3.130017 -0.212954 -0.115804 -1.334983 -0.893730
2  0.330868 -0.065869 -0.401144 -0.144781 -0.382672 -1.334983 -0.509436
3  1.412950 -0.054126 -0.401144 -0.144781 -0.382672 -1.334983 -0.509436
4  0.680463 -0.069939 -2.334096  0.673299 -0.382672 -1.334983  0.028576

      TG      HDL      LDL      VLDL      BMI Gender_M Gender_f \
0 -1.035084  1.810756 -1.085457 -0.369958 -1.124622 False   False
1 -0.678063 -0.158692 -0.457398 -0.342649 -1.326239 True   False
2 -1.035084  1.810756 -1.085457 -0.369958 -1.124622 False   False
3 -1.035084  1.810756 -1.085457 -0.369958 -1.124622 False   False
4 -0.963680 -0.613180 -0.547121 -0.397267 -1.729472 True   False

  CLASS_N CLASS_P CLASS_Y CLASS_Y
0  False   False   False   False
1  False   False   False   False
2  False   False   False   False
3  False   False   False   False
4  False   False   False   False
```

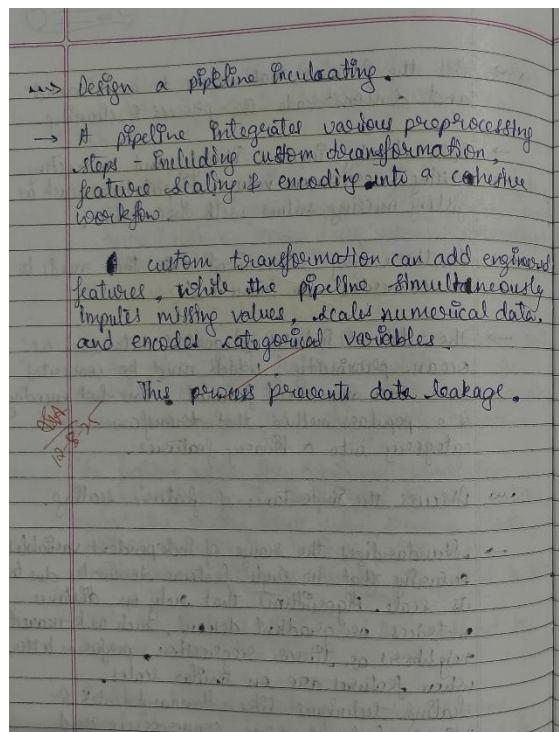
LABORATORY PROGRAM – 2

Demonstrate various data pre-processing techniques for a given dataset

OBSERVATION BOOK

10.3.25	LABORATORY PROGRAM - 2 DEMONSTRATE VARIOUS DATA PRE-PROCESSING TECHNIQUES FOR A GIVEN DATASET	→ Demonstrate the process of creating a test set.
→ Perform the describe and info steps:	→ Random split: <pre>train_set_random, test_set_random = train_test_split(housing['selected'], test_size=0.2, random_state=42)</pre>	
Import pandas as pd. <pre>df = pd.read_csv('housing.csv')</pre> <pre>df.describe()</pre>	→ train_set_random.head() → test_set_random.head()	
→ df.info	→ → Stratified Test set: <pre>split = StratifiedShuffleSplit(n_splits=1, test_size=0.2, random_state=42)</pre> <pre>for train_index, test_index in split.split(housing_selected['income_cat']):</pre> <pre> strat_train_set = housing_selected.loc[train_index]</pre> <pre> strat_train_set = housing_selected.loc[test_index]</pre> <pre> strat_train_set.head()</pre> <pre> strat_train_sets_head()</pre>	

→ List the geographical features from the dataset and plot a graph to visualize geographical data.	→ Data cleaning involves identifying and correcting missing or erroneous values. Imputation such as filling missing values with the median.
→ <pre>housing.plot(kind='scatter', x='longitude', y='latitude', alpha=0.4, s=housing['population']/100, label='population', figsize=(7,5), c='median_house_value', cmap=plt.cm.viridis, colorbar=True)</pre> plt.legend()	→ Is there any categorical data that needs to be converted to numerical?
→ Plot a graph to show features correlation with housing price. Which feature correlates to the maximum. Plot the graph for that with housing price and analyse what the graph indicate.	→ The dataset includes categorical data such as ocean_proximity, which must be converted to a numerical format. One-hot encoding is a popular method that transforms each category into a binary feature.
→ The feature with the maximum correlation is median income. A scatter plot of this feature against the median house value shows a clear positive relationship, indicating that as median incomes increase, housing prices tend to rise.	→ Discuss the importance of feature scaling.
→ List other features that could be combined to improve correlation score.	→ Standardizes the range of independent variables, ensuring that no single feature dominates due to its scale. Algorithms that rely on distance metrics or gradient descent, such as k-nearest neighbors or linear regression, perform better when features are on similar scales.
→ Features such as total_rooms, total_bedrooms, population, and households can be combined to form derived features like income_per_household, bedrooms_per_room, population_per_household. These new features capture underlying patterns.	Scaling techniques like StandardScaler & MinMaxScaler improve convergence and overall model performance.



CODE WITH OUTPUT

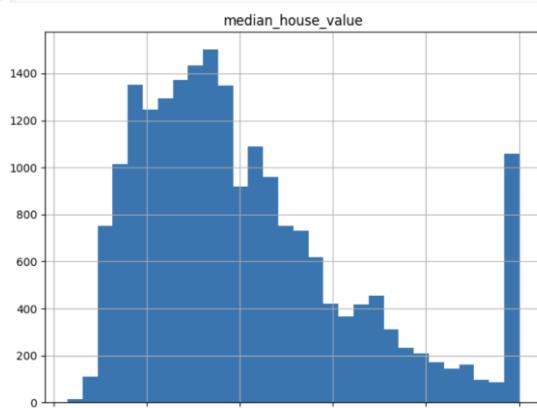
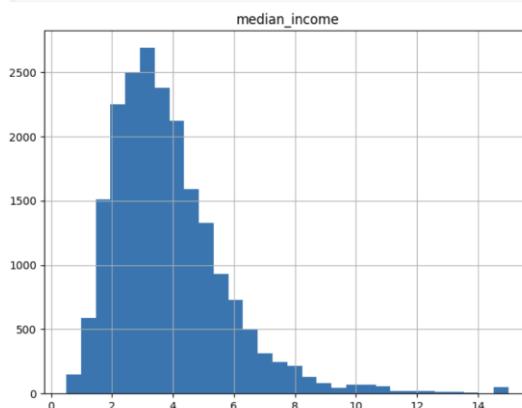
```
# Load the dataset into a pandas DataFrame
df = pd.read_csv('housing.csv')

# Display descriptive statistics
df.describe()
```

	longitude	latitude	housing_median_age	total_rooms	total_bedrooms	population
count	20640.000000	20640.000000	20640.000000	20640.000000	20433.000000	20640.000000
mean	-119.569704	35.631861	28.639486	2635.763081	537.870553	1425.476744
std	2.003532	2.135952	12.585558	2181.615252	421.385070	1132.462122
min	-124.350000	32.540000	1.000000	2.000000	1.000000	3.000000
25%	-121.800000	33.930000	18.000000	1447.750000	296.000000	787.000000
50%	-118.490000	34.260000	29.000000	2127.000000	435.000000	1166.000000
75%	-118.010000	37.710000	37.000000	3148.000000	647.000000	1725.000000
max	-114.310000	41.950000	52.000000	39320.000000	6445.000000	35682.000000

import matplotlib.pyplot as plt
df.hist(column='median_income', bins=30, figsize=(8,6))
plt.show()

import matplotlib.pyplot as plt
df.hist(column='median_house_value', bins=30, figsize=(8,6))
plt.show()



```

import pandas as pd
import numpy as np
from sklearn.model_selection import train_test_split, StratifiedShuffleSplit

# Load the dataset
housing = pd.read_csv('housing.csv')

# For this demonstration, consider only 'median_income' and 'median_house_value'
housing_selected = housing[['median_income', 'median_house_value']].copy()

# Random split: This splits the data randomly without preserving any specific distribution.
train_set_random, test_set_random = train_test_split(housing_selected, test_size=0.2, random_state=42)

# For stratified sampling, first create an income category.
housing_selected['income_cat'] = pd.cut(housing_selected['median_income'],
                                         bins=[0., 1.5, 3.0, 4.5, 6., np.inf],
                                         labels=[1, 2, 3, 4, 5])

# Use StratifiedShuffleSplit to ensure the income distribution is preserved in both sets.
split = StratifiedShuffleSplit(n_splits=1, test_size=0.2, random_state=42)
for train_index, test_index in split.split(housing_selected, housing_selected['income_cat']):
    strat_train_set = housing_selected.loc[train_index]
    strat_test_set = housing_selected.loc[test_index]

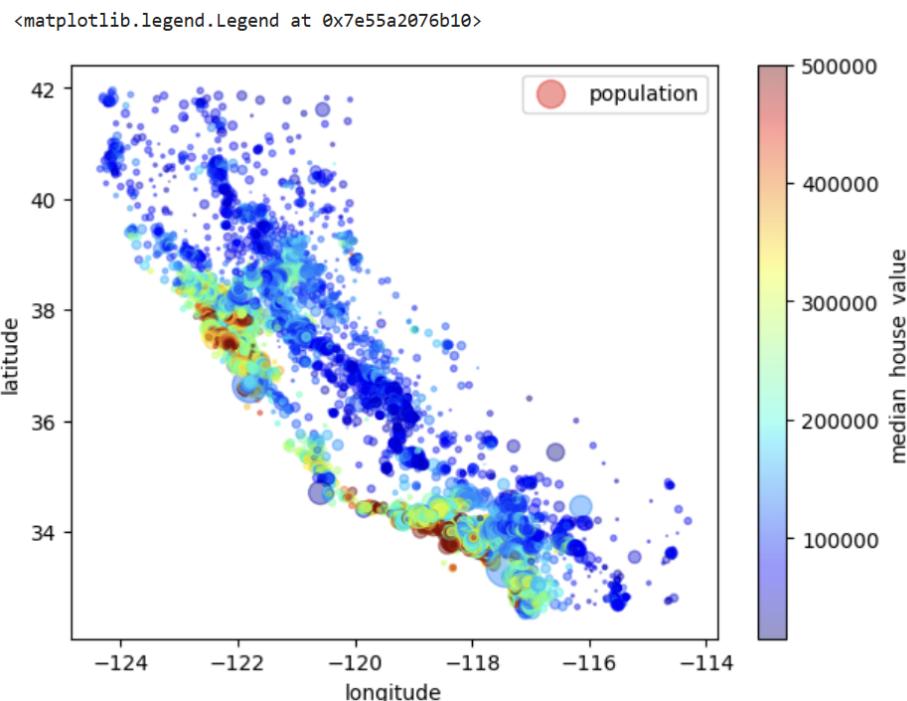
# Remove the temporary income category attribute.
for dataset in (strat_train_set, strat_test_set):
    dataset.drop("income_cat", axis=1, inplace=True)

```

```

import matplotlib.pyplot as plt
housing.plot(kind="scatter", x="longitude", y="latitude", alpha=0.4,
             s=housing["population"]/100, label="population", figsize=(7,5),
             c="median_house_value", cmap=plt.get_cmap("jet"), colorbar=True,)
plt.legend()

```

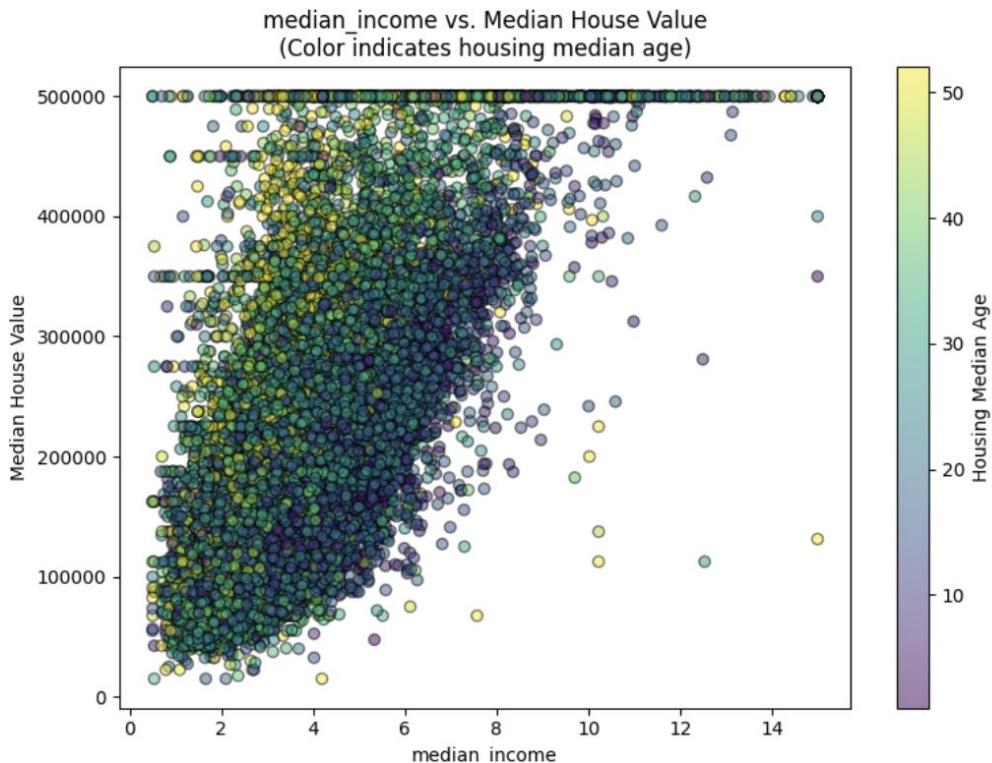


```

import matplotlib.pyplot as plt
import seaborn as sns

plt.figure(figsize=(8,6))
# Differentiate by using 'housing_median_age' for the color
scatter = plt.scatter(housing_numeric[max_feature],
                      housing_numeric["median_house_value"],
                      alpha=0.5,
                      c=housing_numeric["housing_median_age"],
                      cmap='viridis',
                      edgecolor='k')
plt.xlabel(max_feature)
plt.ylabel("Median House Value")
plt.title(f"{max_feature} vs. Median House Value\n(Color indicates housing median age)")
# Add a colorbar to explain the color mapping
cbar = plt.colorbar(scatter)
cbar.set_label("Housing Median Age")
plt.tight_layout()
plt.show()

```



```

from sklearn.preprocessing import OneHotEncoder

# Extract the categorical attribute
housing_cat = housing[["ocean_proximity"]]

# Perform one-hot encoding
encoder = OneHotEncoder()
housing_cat_1hot = encoder.fit_transform(housing_cat).toarray()

# Create a DataFrame for the encoded features
housing_cat_1hot_df = pd.DataFrame(housing_cat_1hot,
                                    columns=encoder.get_feature_names_out(["ocean_proximity"]))
housing_cat_1hot_df.head()
from sklearn.base import BaseEstimator, TransformerMixin
from sklearn.pipeline import Pipeline
from sklearn.compose import ColumnTransformer
from sklearn.preprocessing import StandardScaler

# Custom transformer to add engineered attributes
class CombinedAttributesAdder(BaseEstimator, TransformerMixin):
    def __init__(self, add_bedrooms_per_room=True):

```

```

        self.add_bedrooms_per_room = add_bedrooms_per_room
    def fit(self, X, y=None):
        return self
    def transform(self, X):
        # Assumes X is a NumPy array with the following columns:
        # total_rooms (index 3), total_bedrooms (index 2), population (index 4), households (index 5)
        rooms_per_household = X[:, 3] / X[:, 5]
        population_per_household = X[:, 4] / X[:, 5]
        if self.add_bedrooms_per_room:
            bedrooms_per_room = X[:, 2] / X[:, 3]
            return np.c_[X, rooms_per_household, population_per_household, bedrooms_per_room]
        else:
            return np.c_[X, rooms_per_household, population_per_household]

# Identify numerical and categorical columns
num_attribs = housing.drop("ocean_proximity", axis=1).columns # All numeric columns
cat_attribs = ["ocean_proximity"]

# Build numerical pipeline: impute missing values, add new attributes, then scale
num_pipeline = Pipeline([
    ('imputer', SimpleImputer(strategy="median")),
    ('attribs_adder', CombinedAttributesAdder()),
    ('std_scaler', StandardScaler()),
])
# Build the full pipeline combining numerical and categorical processing
full_pipeline = ColumnTransformer([
    ("num", num_pipeline, num_attribs),
    ("cat", OneHotEncoder(), cat_attribs),
])
# Process the dataset using the pipeline
housing_prepared = full_pipeline.fit_transform(housing)
print("Shape of processed data:", housing_prepared.shape)

```

LABORATORY PROGRAM – 3

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

OBSERVATION BOOK

Date = 17-3-25

LABORATORY PROGRAM -3 X

IMPLEMENT LINEAR AND MULTI-LINEAR
REGRESSION ALGORITHM USING
APPROPRIATE DATASET

SOLVING THE FOLLOWING LINEAR REGRESSION
PROBLEM

$$1. \quad X = \begin{bmatrix} 1 & 1 \\ 1 & 2 \\ 1 & 3 \\ 1 & 4 \end{bmatrix} \quad Y = \begin{bmatrix} 2 \\ 4 \\ 5 \\ 9 \end{bmatrix}$$

$$\hat{B} = ((X^T X)^{-1} X^T) Y$$

$$X^T X = \begin{bmatrix} 4 & 10 \\ 10 & 30 \end{bmatrix} \quad (X^T X)^{-1} = \frac{1}{20} \begin{bmatrix} 30 & -10 \\ -10 & 4 \end{bmatrix} = \begin{bmatrix} 1.5 & -0.5 \\ -0.5 & 0.2 \end{bmatrix}$$

$$X^T Y = \begin{bmatrix} 20 \\ 61 \end{bmatrix}$$

$$\hat{B} = \begin{bmatrix} 1.5 & -0.5 \\ -0.5 & 0.2 \end{bmatrix} \begin{bmatrix} 20 \\ 61 \end{bmatrix} = \begin{bmatrix} -0.5 \\ 2.2 \end{bmatrix} = \begin{bmatrix} B_0 \\ B_1 \end{bmatrix}$$

2. $X = \begin{bmatrix} 1 & 8 \\ 1 & 10 \\ 1 & 12 \end{bmatrix}, Y = \begin{bmatrix} 10 \\ 13 \\ 16 \end{bmatrix}$

$$\hat{B} = (X^T X)^{-1} X^T Y$$

$$X^T X = \begin{bmatrix} 3 & 30 \\ 30 & 308 \end{bmatrix}, (X^T X)^{-1} = \frac{1}{24} \begin{bmatrix} 308 & -30 \\ -30 & 3 \end{bmatrix}$$

$$X^T Y = \begin{bmatrix} 1 & 1 & 1 \\ 8 & 10 & 12 \end{bmatrix} \begin{bmatrix} 10 \\ 13 \\ 16 \end{bmatrix} = \begin{bmatrix} 39 \\ 402 \end{bmatrix}$$

$$\hat{B} = \frac{1}{24} \begin{bmatrix} 308 & -30 \\ -30 & 3 \end{bmatrix} \begin{bmatrix} 39 \\ 402 \end{bmatrix} = \begin{bmatrix} -2 \\ 1.5 \end{bmatrix}$$

The Price for a 20-inch Pizza :

$$y = -2 + 1.5x$$

$$y = -2 + 1.5 \times 20 = -2 + 30 = 28$$

\therefore The Predicted price is \$28

After building the regression model,

1. Data Preprocessing steps 1-5 checked for missing values and imputed numeric columns with their mean. Textual numbers were converted to numeric, and categorical variables (like 'State') were one-hot encoded. Scaling was considered when feature ranges differed significantly.
2. Visualization for Canada per capita income: I plotted the regression line with the data points, which showed a clear upward trend. This indicates that per capita income increases over the years. The linear trend confirms that linear regression is a suitable model.
3. Predicted Salary for Hiring Data: The prediction for a candidate with 15 years of experience, a test score of 10, and an interview score of 10 is \$86,244.67.
4. Preprocessing for 1000 Companies.csv: I encoded the 'State' column using one-hot encoding (pd.get_dummies with drop first=True) to avoid multicollinearity. Scaling was considered for features like R&D Spend, Admin -ilation and Marketing Spend due to their different ranges to enhance model stability.

CODE WITH OUTPUT

```
import pandas as pd
from sklearn.model_selection import train_test_split
from sklearn.tree import DecisionTreeClassifier
from sklearn.metrics import accuracy_score, confusion_matrix, classification_report
import matplotlib.pyplot as plt
from sklearn.tree import plot_tree

# Load the iris dataset (make sure iris.csv is in the working directory)
iris = pd.read_csv("iris.csv")
# Assuming the last column is the target (species) and the rest are features.
X = iris.iloc[:, :-1]
y = iris.iloc[:, -1]

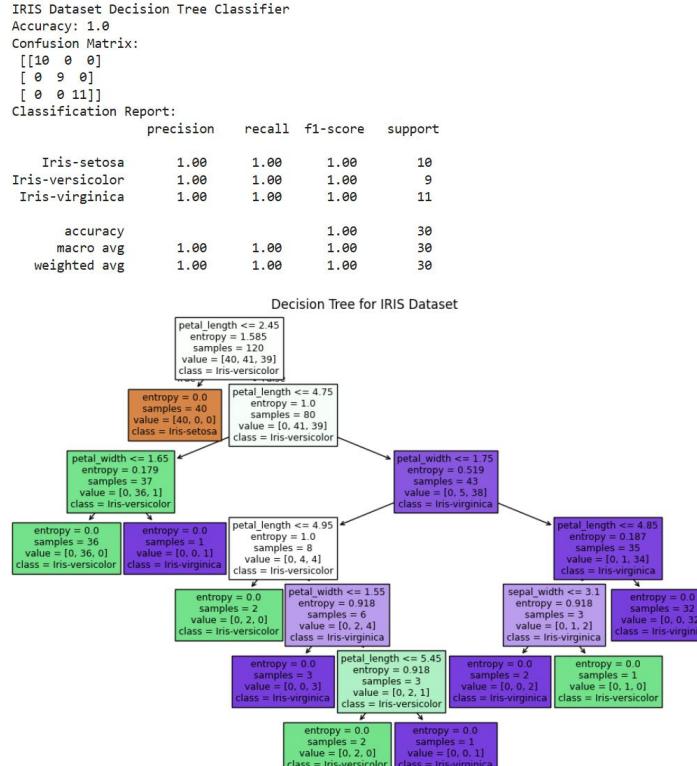
# Split data into training and testing sets (80% training, 20% testing)
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)

# Initialize and train the Decision Tree classifier
clf_iris = DecisionTreeClassifier(criterion='entropy', random_state=42)
clf_iris.fit(X_train, y_train)

# Make predictions and evaluate the model
y_pred_iris = clf_iris.predict(X_test)
accuracy_iris = accuracy_score(y_test, y_pred_iris)
conf_matrix_iris = confusion_matrix(y_test, y_pred_iris)

print("IRIS Dataset Decision Tree Classifier")
print("Accuracy:", accuracy_iris)
print("Confusion Matrix:\n", conf_matrix_iris)
print("Classification Report:\n", classification_report(y_test, y_pred_iris))

# Visualize the decision tree
plt.figure(figsize=(12, 8))
plot_tree(clf_iris, filled=True, feature_names=X.columns, class_names=clf_iris.classes_)
plt.title("Decision Tree for IRIS Dataset")
plt.show()
```



```

import pandas as pd
from sklearn.model_selection import train_test_split
from sklearn.tree import DecisionTreeClassifier
from sklearn.metrics import accuracy_score, confusion_matrix, classification_report
import matplotlib.pyplot as plt
from sklearn.tree import plot_tree

# Load the drug dataset (make sure drug.csv is in the working directory)
drug = pd.read_csv("drug.csv")

# Since the target column is 'Drug', drop it from the features
X_drug = drug.drop('Drug', axis=1)
y_drug = drug['Drug']

# If there are categorical features, perform necessary encoding
from sklearn.preprocessing import LabelEncoder
le = LabelEncoder()
# Encode features that are categorical
for col in X_drug.select_dtypes(include='object').columns:
    X_drug[col] = le.fit_transform(X_drug[col])
# Also encode the target variable if necessary
y_drug = le.fit_transform(y_drug)

# Split the data (80% training, 20% testing)
X_train_d, X_test_d, y_train_d, y_test_d = train_test_split(X_drug, y_drug, test_size=0.2, random_state=42)

# Initialize and train the Decision Tree classifier using entropy criterion
clf_drug = DecisionTreeClassifier(criterion='entropy', random_state=42)
clf_drug.fit(X_train_d, y_train_d)

# Make predictions and evaluate the model
y_pred_drug = clf_drug.predict(X_test_d)
accuracy_drug = accuracy_score(y_test_d, y_pred_drug)
conf_matrix_drug = confusion_matrix(y_test_d, y_pred_drug)

print("Drug Dataset Decision Tree Classifier")
print("Accuracy: ", accuracy_drug)
print("Confusion Matrix:\n", conf_matrix_drug)
print("Classification Report:\n", classification_report(y_test_d, y_pred_drug))

# Visualize the decision tree
plt.figure(figsize=(12, 8))
plot_tree(clf_drug, filled=True, feature_names=X_drug.columns,
          class_names=[str(cls) for cls in clf_drug.classes_])
plt.title("Decision Tree for Drug Dataset")
plt.show()

```

Drug Dataset Decision Tree Classifier

Accuracy: 1.0

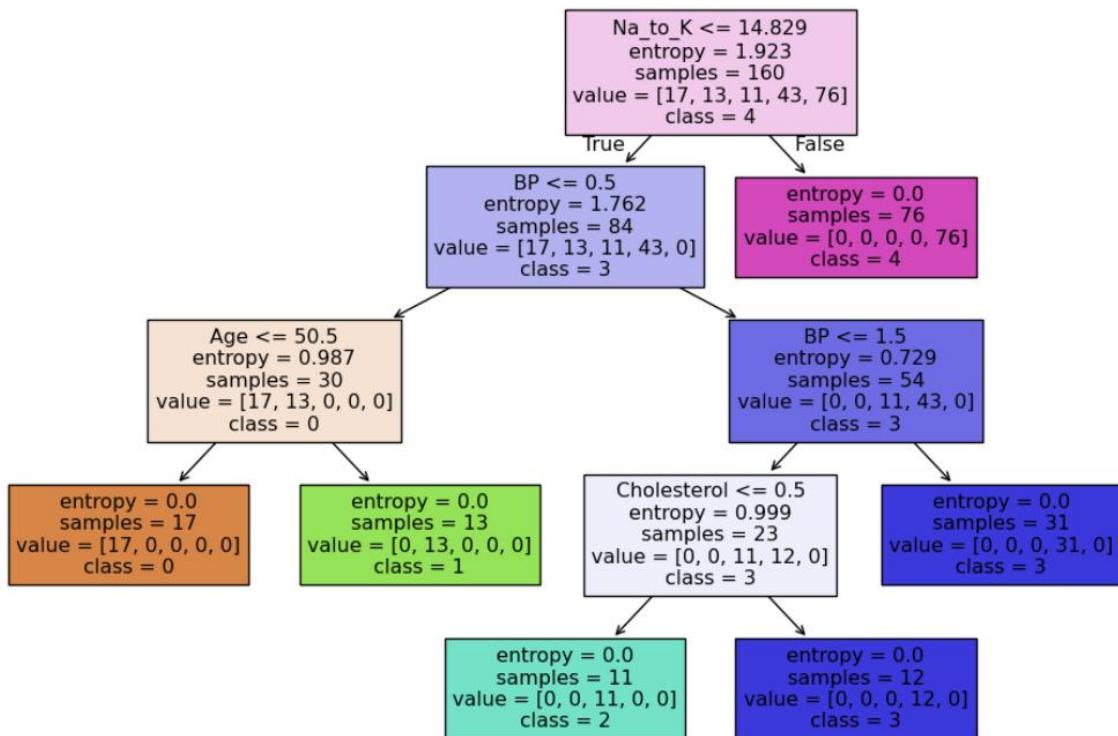
Confusion Matrix:

```
[[ 6  0  0  0  0]
 [ 0  3  0  0  0]
 [ 0  0  5  0  0]
 [ 0  0  0 11  0]
 [ 0  0  0  0 15]]
```

Classification Report:

	precision	recall	f1-score	support
0	1.00	1.00	1.00	6
1	1.00	1.00	1.00	3
2	1.00	1.00	1.00	5
3	1.00	1.00	1.00	11
4	1.00	1.00	1.00	15
accuracy			1.00	40
macro avg	1.00	1.00	1.00	40
weighted avg	1.00	1.00	1.00	40

Decision Tree for Drug Dataset



LABORATORY PROGRAM – 4

Implement Linear and Multi-Linear Regression algorithm using appropriate dataset

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LABORATORY PROGRAM - 4

Use an appropriate dataset for building the decision tree (IRIS) and apply this knowledge to classify a new sample.

o Considering the dataset, calculate entropy and Information gain w.r.t. to target variable Classification.

→ Total instances : 5

o Classification counts :

- No : 4 (Instances : 1, 2, 6, 7)
- Yes : 1 (Instance : 8)

$$E(S) = -P_1 \log_2 P_1 - P_2 \log_2 P_2$$

$$P_1 = \frac{4}{5} \rightarrow P_1 = \frac{1}{5}$$

$$E(S) = -(0.8 \log_2 0.8 + 0.2 \log_2 0.2)$$

$$E(S) = -(0.8 \times (-0.32193)) + 0.2 \times (-2.32193)$$

$$E(S) = -0.72193$$

→ Information gain for Attribute a_2 :

Value : Hot, Cool

→ For $a_2 = \text{Hot}$:

Instances : 1, 2, 7, 8

Classification : No, No, No, Yes

$$E(\text{Hot}) = -\left(\frac{3}{4} \log_2 \frac{3}{4} + \frac{1}{4} \log_2 \frac{1}{4}\right)$$

$$E(\text{Hot}) = -(0.75 \times -0.4150 + 0.25 \times -2)$$

$$E(\text{Hot}) = 0.81125$$

→ For $a_2 = \text{Cool}$:

Instances : 6

Classification : No, Entropy for $a_2 = \text{Cool}$ is 0

Weighted Entropy for a_2 :

$$E(S, a_2) = \frac{4}{5} \times 0.81125 + \frac{1}{5} \times 0 = 0.649$$

Information gain for a_2 :

$$IG(a_2) = E(S) - E(S, a_2) = 0.72193 - 0.649$$

$$\rightarrow 0.07293$$

o After building the Decision Tree Model:

1. For "iris.csv" dataset:
- o The model achieved an accuracy of 1.00 (100%) on the IRIS dataset
- o Confusion Matrix Interpretation:
 - ↳ Each diagonal entry represents the no. of correctly classified instances for each class (Bio-logical, Bio-chemical, Bio-medical).
 - ↳ Since all off-diagonal elements are zero, there were no misclassifications.
 - ↳ This indicates that the model perfectly distinguished b/w the 3 classes without confusing any of them.
2. For "petrol consumption.csv" dataset:
- o Regression Tree Structure:
 - ↳ Split data into regions based on features.
 - ↳ Each leaf node predicts a continuous Petrol Consumption value.
 - ↳ Features near the root indicate the most important predictors.
- o Key Evaluation Metrics:
 - MAE : 94.3
 - MSE : 17347.7
 - RMSE : 131.71

Handling Continuous Targets vs Categorical Labels	
→ Regression Tree:	
↳ Predict continuous outcomes	
↳ Split chosen to minimize variance / MSE	
↳ Leaf nodes output mean target values	
→ Decision Tree Classifier:	
↳ Predict discrete class labels	
↳ Split chosen to maximize node purity	
↳ Leaf nodes assign the majority class labels	

CODE WITH OUTPUT

```

import pandas as pd
from sklearn.linear_model import LinearRegression
# Load the data
income_data = pd.read_csv("canada_per_capita_income.csv")
# Assumed data columns: 'Year' and 'PerCapitaIncome'
print("Canada Income Data Head:")
print(income_data.head())
# Prepare feature and target
X_income = income_data[["year"]]    # Predictor variable: Year
y_income = income_data["per capita income (US$)"]
# Build and train the linear regression model
model_income = LinearRegression()
model_income.fit(X_income, y_income)

# Predict per capita income for the year 2020
predicted_income = model_income.predict([[2020]])

print("\nPredicted per capita income for Canada in 2020:", predicted_income[0])

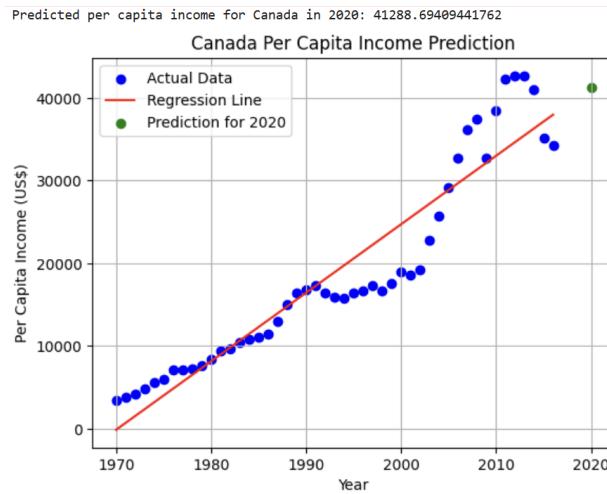
# Plot the data points and the regression line
plt.scatter(X_income, y_income, color='blue', label='Actual Data')
plt.plot(X_income, model_income.predict(X_income), color='red', label='Regression Line')

# Plot the prediction for 2020
plt.scatter(2020, predicted_income[0], color='green', label='Prediction for 2020')

# Customize the plot
plt.xlabel('Year')
plt.ylabel('Per Capita Income (US$)')
plt.title('Canada Per Capita Income Prediction')
plt.legend()
plt.grid(True)

# Display the plot
plt.show()

```



```

import numpy as np
import matplotlib.pyplot as plt
import pandas as pd
from sklearn.linear_model import LinearRegression

# Load the salary data
salary_data = pd.read_csv("salary.csv")
print(salary_data.head())

# Prepare feature and target
X_salary = salary_data[["YearsExperience"]] # Predictor variable: Years of Experience
y_salary = salary_data["Salary"]

# Build and train the linear regression model
model_salary = LinearRegression()
model_salary.fit(X_salary, y_salary)

import matplotlib.pyplot as plt
# Plot the data points and the regression line
plt.scatter(X_salary, y_salary, color='blue', label='Actual Data')
plt.plot(X_salary, model_salary.predict(X_salary), color='red', label='Regression Line')

# Plot the prediction for 12 years of experience
plt.scatter(12, predicted_salary[0], color='green', label='Prediction for 12 years')

# Customize the plot
plt.xlabel('Years of Experience')
plt.ylabel('Salary')
plt.title('Salary Prediction based on Experience')
plt.legend()
plt.grid(True)

# Display the plot
plt.show()

```

Predicted salary for an employee with 12 years of experience: 139980.88923969213



```

import pandas as pd
import numpy as np
from sklearn.linear_model import LinearRegression

# Read the CSV file (ensure the file is uploaded in your Colab environment)
df = pd.read_csv("hiring.csv")

# Rename columns for convenience
df.columns = ['experience', 'test_score', 'interview_score', 'salary']

print("Original Data:")
print(df)
# Function to convert experience values to numeric
def convert_experience(x):
    try:
        return float(x)
    except:
        x_lower = str(x).strip().lower()
        return num_map.get(x_lower, np.nan)

# Convert the 'experience' column using the mapping
df['experience'] = df['experience'].apply(convert_experience)

# Convert 'test_score', 'interview_score', and 'salary' to numeric (coerce errors to NaN)
df['test_score'] = pd.to_numeric(df['test_score'], errors='coerce')
df['interview_score'] = pd.to_numeric(df['interview_score'], errors='coerce')
df['salary'] = pd.to_numeric(df['salary'], errors='coerce')

print("\nData After Conversion:")
print(df)

# Fill missing values in numeric columns using the column mean
df['experience'].fillna(df['experience'].mean(), inplace=True)
df['test_score'].fillna(df['test_score'].mean(), inplace=True)
df['interview_score'].fillna(df['interview_score'].mean(), inplace=True)

print("\nData After Filling Missing Values:")
print(df)

```

```

# Prepare the feature matrix X and target vector y
X = df[['experience', 'test_score', 'interview_score']]
y = df['salary']

# Build and train the Multiple Linear Regression model
model = LinearRegression()
model.fit(X, y)

# Predict salaries for the given candidate profiles
# Candidate 1: 2 years of experience, 9 test score, 6 interview score
candidate1 = np.array([[2, 9, 6]])
predicted_salary1 = model.predict(candidate1)

# Candidate 2: 12 years of experience, 10 test score, 10 interview score
candidate2 = np.array([[12, 10, 10]])
predicted_salary2 = model.predict(candidate2)

print("\nPredicted Salary for Candidate (2 yrs, 9 test, 6 interview): $", round(predicted_salary1[0], 2))
print("Predicted Salary for Candidate (12 yrs, 10 test, 10 interview): $", round(predicted_salary2[0], 2))

import matplotlib.pyplot as plt

# Create the plot
plt.figure(figsize=(10, 6)) # Adjust figure size for better visualization
plt.scatter(df['experience'], y, color='blue', label='Actual Salary') # Plot actual salary against years of experience

# Plot the regression line (this is an approximation since it's a multi-variable regression)
# You can visualize a single feature against the predicted salary
plt.plot(df['experience'], model.predict(X), color='red', label='Regression Line')

# Highlight predictions
plt.scatter(candidate1[0, 0], predicted_salary1, color='green', label='Candidate 1 Prediction')
plt.scatter(candidate2[0, 0], predicted_salary2, color='purple', label='Candidate 2 Prediction')

# Add labels and title
plt.xlabel("Years of Experience")
plt.ylabel("Salary")
plt.title("Salary Prediction based on Experience, Test Score, Interview Score")

# Add a legend
plt.legend()
plt.grid(True)
plt.show()

```



LABORATORY PROGRAM – 5

Build Logistic Regression Model for a given dataset

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LABORATORY PROGRAM - 5

Built logistic regression Model for a given dataset

ws Binary Classification : Predicting Student Pass/Fail :

a) Logistic Regression Equation :

$$Z = a_0 + a_1 x$$

$$P(y=1) = \frac{1}{1 + e^{-(a_0 + a_1 x)}}$$

$$a_0 = -5 \text{ and } a_1 = 0.8$$

$$P(\text{Pass}) = \frac{1}{1 + e^{-(5+0.8x)}}$$

b) Calculate the probability for 7 Study Hours

$$P(\text{Pass}) = \frac{1}{1 + e^{-(5+0.8 \times 7)}} = \frac{1}{e^{-0.6} + 1}$$

$$P(\text{Pass}) = 0.646$$

Thus the probability that the student will pass is approx 64.6%.

c) Predicted Class Based on a Threshold of 0.5

$$P(\text{Pass}) \geq 0.5 \Rightarrow \text{Pass}$$

$$P(\text{Pass}) < 0.5 \Rightarrow \text{Fail}$$

$$\therefore P(\text{Pass}) = 0.646 \Rightarrow \text{Pass}$$

ws Multi Class Classification: Applying the Softmax function:

$$z = [2, 1, 0], P_i = \frac{e^{z_i}}{\sum e^{z_j}}$$

$$e^2 = 7.389, e^1 = 2.718, e^0 = 1$$

$$\sum e^{z_j} = 11.107$$

The Softmax Probabilities:

$$P_1 = \frac{7.389}{11.107} \approx 0.665$$

$$P_2 = \frac{2.718}{11.107} \approx 0.245$$

$$P_3 = \frac{1}{11.107} \approx 0.090$$

The Softmax probability values for 3 classes:
Class 1: 66.5%, Class 2: 24.5%, Class 3: 9.0%

ws After building the logistic regression model for dataset file "HR comma sep.csv":
 Variable include satisfaction level, time spent at the company, no. of projects, average monthly hours, salary, work accidents history and promotions have impact on employee retention.
 Employee with low satisfaction, working more hours, low salary, or no promotions are more likely to leave.
 The logistic regression model achieved 76% accuracy, which is decent but can be improved. Advanced models like random forests or neural networks could better capture complex employee behaviour patterns.

CODE WITH OUTPUT

```
import pandas as pd
from matplotlib import pyplot as plt
# %matplotlib inline
#"%matplotlib inline" will make your plot outputs appear and be stored within the notebook.

df = pd.read_csv("insurance_data.csv")
df.head()

plt.scatter(df.age,df.bought_insurance,marker='+',color='red')

from sklearn.model_selection import train_test_split
X_train, X_test, y_train, y_test = train_test_split(df[['age']], df.bought_insurance, train_size=0.9, random_state=10)
X_train.shape

X_test

from sklearn.linear_model import LogisticRegression
model = LogisticRegression()

model.fit(X_train, y_train)

X_test

y_test

y_predicted = model.predict(X_test)
y_predicted

model.score(X_test,y_test)

model.predict_proba(X_test)

y_predicted = model.predict([[60]])
y_predicted

#model.coef_ indicates value of m in y=m*x + b equation
model.coef_

#model.intercept_ indicates value of b in y=m*x + b equation
model.intercept_

#Lets defined sigmoid function now and do the math with hand
import math
def sigmoid(x):
    return 1 / (1 + math.exp(-x))

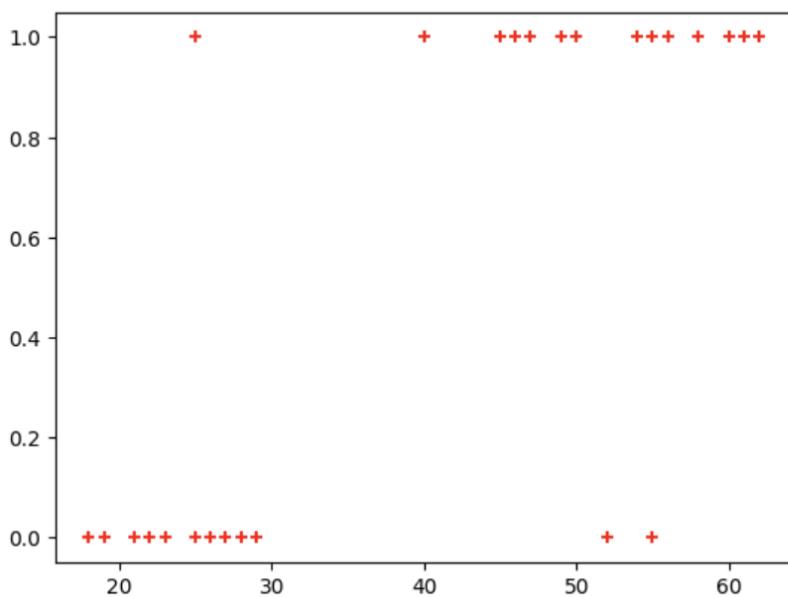
def prediction_function(age):
    z = 0.127 * age - 4.973 # 0.12740563 ~ 0.0127 and -4.97335111 ~ -4.97
    y = sigmoid(z)
    return y

age = 35
prediction_function(age)

"""0.37 is less than 0.5 which means person with 35 will not buy the insurance"""


```

'0.37 is less than 0.5 which means person with 35 will not buy the insurance'



```
# Import necessary libraries
import pandas as pd
from sklearn.datasets import load_iris
from sklearn.model_selection import train_test_split
from sklearn.linear_model import LogisticRegression
from sklearn.metrics import accuracy_score
from sklearn import metrics
import matplotlib.pyplot as plt

# Load the Iris dataset
iris = pd.read_csv("iris.csv")
iris.head()

X=iris.drop('species',axis='columns')# Features (sepal length, sepal width, petal length, petal width)
y = iris.species # Target labels (0: Setosa, 1: Versicolor, 2: Virginica)

# Split the dataset into 80% training and 20% testing
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)

# Initialize the Multinomial Logistic Regression model
# Use 'multinomial' for multi-class classification and 'lbfgs' solver
model = LogisticRegression(multi_class='multinomial')

# Train the model on the training data
model.fit(X_train, y_train)

# Make predictions on the test data
y_pred = model.predict(X_test)

# Calculate the accuracy of the model on the test data
accuracy = accuracy_score(y_test, y_pred)

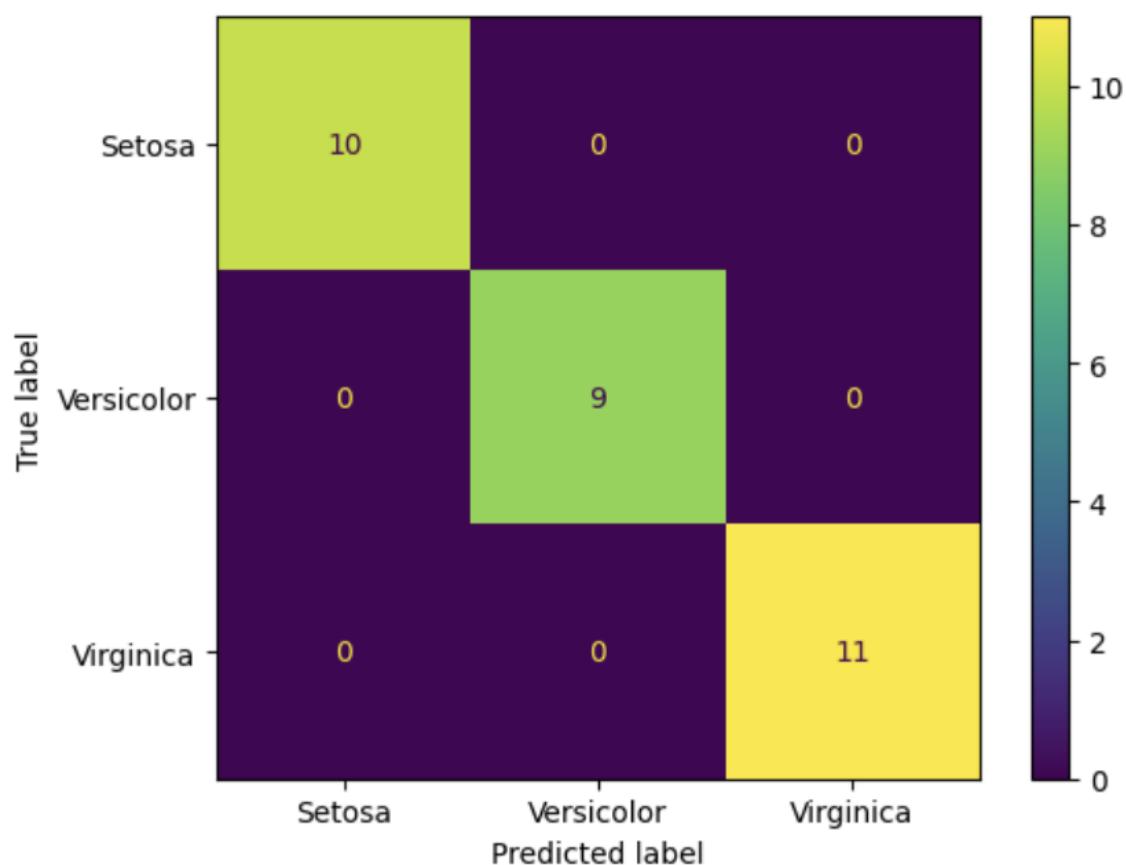
# Display the accuracy
print(f"Accuracy of the Multinomial Logistic Regression model on the test set: {accuracy:.2f}")

confusion_matrix = metrics.confusion_matrix(y_test, y_pred)

cm_display = metrics.ConfusionMatrixDisplay(confusion_matrix = confusion_matrix, display_labels = ["Setosa",
"Versicolor", "Virginica"])

cm_display.plot()
plt.show()
```

Accuracy of the Multinomial Logistic Regression model on the test set: 1.00



LABORATORY PROGRAM – 6

Build KNN Classification model for a given dataset.

OBSERVATION BOOK

LABORATORY PROGRAM - 6																																																					
Build KNN Classification Model for A Given Dataset																																																					
Considering the dataset, for K=3 and test data (x, 35, 100) as (Person, Age, SalaryK) solve using KNN classifier Model and Predict its target.																																																					
<table border="1"><thead><tr><th>Person</th><th>Age</th><th>SalaryK</th><th>Target</th><th>Distance</th><th>Rank</th></tr></thead><tbody><tr><td>A</td><td>18</td><td>50</td><td>N</td><td>52.82</td><td>5</td></tr><tr><td>B</td><td>23</td><td>55</td><td>N</td><td>46.57</td><td>4</td></tr><tr><td>C</td><td>24</td><td>70</td><td>N</td><td>31.96</td><td>2</td></tr><tr><td>D</td><td>21</td><td>60</td><td>Y</td><td>40.45</td><td>3</td></tr><tr><td>E</td><td>43</td><td>70</td><td>Y</td><td>31.04</td><td>1</td></tr><tr><td>F</td><td>38</td><td>40</td><td>Y</td><td>60.08</td><td>6</td></tr><tr><td>X</td><td>35</td><td>100</td><td>?</td><td>?</td><td>?</td></tr></tbody></table>						Person	Age	SalaryK	Target	Distance	Rank	A	18	50	N	52.82	5	B	23	55	N	46.57	4	C	24	70	N	31.96	2	D	21	60	Y	40.45	3	E	43	70	Y	31.04	1	F	38	40	Y	60.08	6	X	35	100	?	?	?
Person	Age	SalaryK	Target	Distance	Rank																																																
A	18	50	N	52.82	5																																																
B	23	55	N	46.57	4																																																
C	24	70	N	31.96	2																																																
D	21	60	Y	40.45	3																																																
E	43	70	Y	31.04	1																																																
F	38	40	Y	60.08	6																																																
X	35	100	?	?	?																																																
The 3 Nearest Neighbors to X are E : With target Y C : With Target N D : With Target Y																																																					
Thus, by majority vote; the predicted target for X is Y. That is ; (x, 35, 100) has target 'Y'.																																																					
1. For Iris Dataset: How to choose the K value? Demonstrate using accuracy rate and error rate. Experiment with different k values and observe the accuracy and error rates. For instance with k=3 we achieved a accuracy of 1.0(100%) and an error rate of 0 (Since error = 0 - accuracy) 2. Demonstrate using accuracy.																																																					
2. For Diabetes data set What is the purpose of feature scaling? How to perform it? Since KNN uses distance measure, features with larger ranges can dominate the calculation. Scaling ensures that each feature contributes equally.																																																					
We used scikit-learn's StandardScaler to standardize features by subtracting the mean & dividing by the standard deviation, leading to improved model performance.																																																					

CODE WITH OUTPUT

```
# Import necessary libraries
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns

# For model building and evaluation
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import StandardScaler
from sklearn.neighbors import KNeighborsClassifier
from sklearn.metrics import accuracy_score, confusion_matrix, classification_report

# ----- Part 1: IRIS Dataset ----- #
# Load the iris dataset (ensure iris.csv is in the same directory or provide correct path)
iris_df = pd.read_csv("iris.csv")

# Separate features and target
X_iris = iris_df.drop("species", axis=1)
y_iris = iris_df["species"]

# Split the data (80% training, 20% testing)
X_train_iris, X_test_iris, y_train_iris, y_test_iris = train_test_split(
    X_iris, y_iris, test_size=0.2, random_state=42
)

# Choose a value for k; here K=3 is used as an example.
knn_iris = KNeighborsClassifier(n_neighbors=3)
```

```

# Train the model on training data
knn_iris.fit(X_train_iris, y_train_iris)

# Predict on test data
y_pred_iris = knn_iris.predict(X_test_iris)

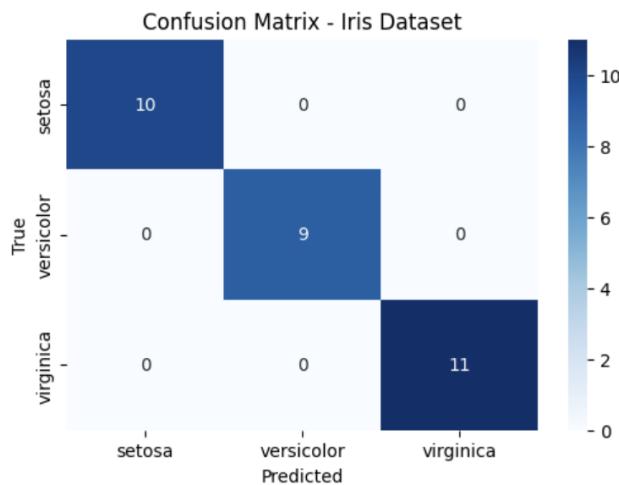
# Calculate accuracy score
acc_iris = accuracy_score(y_test_iris, y_pred_iris)
print("IRIS Dataset Accuracy Score:", acc_iris)

# Compute confusion matrix and classification report
cm_iris = confusion_matrix(y_test_iris, y_pred_iris)
print("\nIRIS Dataset Confusion Matrix:\n", cm_iris)

cr_iris = classification_report(y_test_iris, y_pred_iris)
print("\nIRIS Dataset Classification Report:\n", cr_iris)

```

	precision	recall	f1-score	support
setosa	1.00	1.00	1.00	10
versicolor	1.00	1.00	1.00	9
virginica	1.00	1.00	1.00	11
accuracy			1.00	30
macro avg	1.00	1.00	1.00	30
weighted avg	1.00	1.00	1.00	30



```

# ----- Part 2: Diabetes Dataset ----- #
# Load the diabetes dataset (ensure diabetes.csv is in the same directory or provide correct path)
diabetes_df = pd.read_csv("diabetes.csv")

# Separate features and target (Outcome column is assumed to be the target)
X_diabetes = diabetes_df.drop("Outcome", axis=1)
y_diabetes = diabetes_df["Outcome"]

# Perform feature scaling on the features
scaler = StandardScaler()
X_scaled_diabetes = scaler.fit_transform(X_diabetes)

# Split the scaled data (80% training, 20% testing)
X_train_diab, X_test_diab, y_train_diab, y_test_diab = train_test_split(
    X_scaled_diabetes, y_diabetes, test_size=0.2, random_state=42
)

# Choose a value for k; here K=5 is used as an example.
knn_diabetes = KNeighborsClassifier(n_neighbors=5)

# Train the model on training data

```

```

knn_diabetes.fit(X_train_diab, y_train_diab)

# Predict on test data
y_pred_diab = knn_diabetes.predict(X_test_diab)

# Calculate accuracy score
acc_diab = accuracy_score(y_test_diab, y_pred_diab)
print("Diabetes Dataset Accuracy Score:", acc_diab)

# Compute confusion matrix and classification report
cm_diab = confusion_matrix(y_test_diab, y_pred_diab)
print("\nDiabetes Dataset Confusion Matrix:\n", cm_diab)

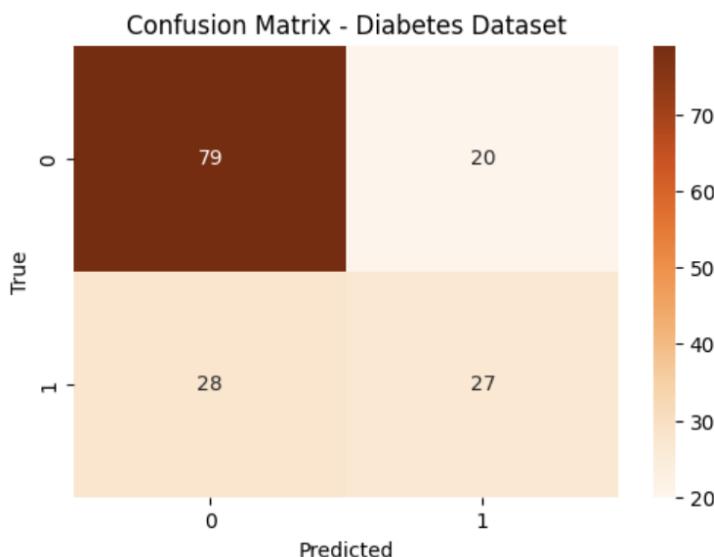
```

```

cr_diab = classification_report(y_test_diab, y_pred_diab)
print("\nDiabetes Dataset Classification Report:\n", cr_diab)

```

	precision	recall	f1-score	support
0	0.74	0.80	0.77	99
1	0.57	0.49	0.53	55
accuracy			0.69	154
macro avg	0.66	0.64	0.65	154
weighted avg	0.68	0.69	0.68	154



```

# ----- Load the Dataset ----- #
# Load heart.csv (make sure the file is in your working directory)
heart_df = pd.read_csv("heart.csv")

# Display the first few rows to check the data
heart_df.head()

# ----- Data Preparation ----- #
# Separate features and target
X_heart = heart_df.drop("target", axis=1)
y_heart = heart_df["target"]

# Perform feature scaling (important for distance-based algorithms like KNN)
scaler = StandardScaler()
X_scaled = scaler.fit_transform(X_heart)

# Split data into training and testing sets (80% train, 20% test)

```

```

X_train, X_test, y_train, y_test = train_test_split(X_scaled, y_heart, test_size=0.2, random_state=42)
# ----- Finding the Best k -----
# We will try a range of k values (neighbors) and select the one with maximum accuracy.
k_range = range(1, 21)
accuracy_scores = []

for k in k_range:
    knn = KNeighborsClassifier(n_neighbors=k)
    knn.fit(X_train, y_train)
    y_pred = knn.predict(X_test)
    acc = accuracy_score(y_test, y_pred)
    accuracy_scores.append(acc)
    print(f"K = {k} --> Accuracy: {acc:.4f}")

        K = 1 --> Accuracy: 0.8525
        K = 2 --> Accuracy: 0.8197
        K = 3 --> Accuracy: 0.8689
        K = 4 --> Accuracy: 0.8852
        K = 5 --> Accuracy: 0.9180
        K = 6 --> Accuracy: 0.9344
        K = 7 --> Accuracy: 0.9180
        K = 8 --> Accuracy: 0.8525
        K = 9 --> Accuracy: 0.8852
        K = 10 --> Accuracy: 0.8852
        K = 11 --> Accuracy: 0.8852
        K = 12 --> Accuracy: 0.8689
        K = 13 --> Accuracy: 0.8852
        K = 14 --> Accuracy: 0.8689
        K = 15 --> Accuracy: 0.9016
        K = 16 --> Accuracy: 0.8852
        K = 17 --> Accuracy: 0.8852
        K = 18 --> Accuracy: 0.9016
        K = 19 --> Accuracy: 0.8852
        K = 20 --> Accuracy: 0.8852

|: # Determine the best k value
best_k = k_range[np.argmax(accuracy_scores)]
print("\nBest k value:", best_k)

Best k value: 6

# ----- Train Final Model with Best k -----
best_knn = KNeighborsClassifier(n_neighbors=best_k)
best_knn.fit(X_train, y_train)
y_pred_best = best_knn.predict(X_test)

# Compute final accuracy, confusion matrix and classification report
final_accuracy = accuracy_score(y_test, y_pred_best)
cm = confusion_matrix(y_test, y_pred_best)
cr_text = classification_report(y_test, y_pred_best)
print("\nFinal Accuracy Score:", final_accuracy)
print("\nConfusion Matrix:\n", cm)
print("\nClassification Report:\n", cr_text)

Final Accuracy Score: 0.9344262295081968

Confusion Matrix:
[[28  1]
 [ 3 29]]

Classification Report:
      precision    recall  f1-score   support
          0       0.90      0.97      0.93      29
          1       0.97      0.91      0.94      32

      accuracy                           0.93      61
     macro avg       0.93      0.94      0.93      61
  weighted avg       0.94      0.93      0.93      61

```

LABORATORY PROGRAM – 7

Build Support vector machine model for a given dataset

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LABORATORY PROGRAM - 7

BUILT SUPPORT VECTOR MACHINE FOR A GIVEN DATASET

Given points $(1, 1), (1, -1)$ and $(6, 0)$ belong to positive class and points $(1, 0), (0, 1)$ & $(0, -1)$ belong to negative class. Draw an optimal hyperplane.

- Point-to-line distance: $Ax + By + C = 0$ and point (x_0, y_0) $\frac{|Ax_0 + By_0 + C|}{\sqrt{A^2 + B^2}}$
- Derivation for these points:

 - Positive: $(1, 1), (1, -1), (6, 0)$
 - Negative: $(1, 0), (0, 1), (0, -1)$

- Shape of separating line: $x = b$, because the points differ only in their x value.
- Nearest positive support points are $(1, 1), (1, -1)$. Thus: $x = b$ is $|x - b| = |1 - b|$.
- Nearest negative support point is $(0, 0)$. Thus: $x = b$ is $|x - b| = |b|$.

Maximizing the minimal distance:

$$b - 1 = 1 - b \Rightarrow 2b = 2 \Rightarrow b = 1$$

Hence margin on each is $1 - 1 = 1$.

Final Hyperplane and Margin Line:

Decision boundary: $x = 1$; $w = [1, 0]$, $b = 1$
 $x + b = 0$

Margin-defining lines: $x = 1$ and $x = -1$
These pass through the support vectors $(1, 1)$ and $(1, -1)$ respectively.

OBSERVATION:

1. For "iris.csv" dataset:

- Accuracy: Linear kernel: 100%
RBF Kernel: 100/100
- Radial Basis Function Kernel gave better performance.
RBF Kernel is capable of capturing non-linear relationships in data by mapping input features into a higher-dimensional space.
- For "Letter-Recognition.csv" dataset:

 - Confusion Matrix Interpretation: The SVM model achieved an accuracy of 95% on the test data. The confusion matrix shows how each letter was predicted versus its actual class.
 - AUC Score: The micro-average AUC score is 1.00, which means the model is very effective at ranking predictions - it assigns higher scores to correct classes with high confidence.

Performance Comparison: SVM on IRIS vs Letter-Recognition Dataset:

Aspect	IRIS Dataset	Letter Recognition Dataset
Classification rate	95%	95%
Accuracy	100/100	95%
AUC	Not computed	10 (Micro Avg)

CODE WITH OUTPUT

```

import numpy as np
import matplotlib.pyplot as plt
from sklearn.svm import SVC

# Data points
X = np.array([[4, 1], [4, -1], [6, 0], [1, 0], [0, 1], [0, -1]])
y = np.array([1, 1, 1, -1, -1, -1])

# Fit linear SVM with a very large C to approximate hard-margin
clf = SVC(kernel='linear', C=1e6)
clf.fit(X, y)

# Extract model parameters
w = clf.coef_[0]
b = clf.intercept_[0]

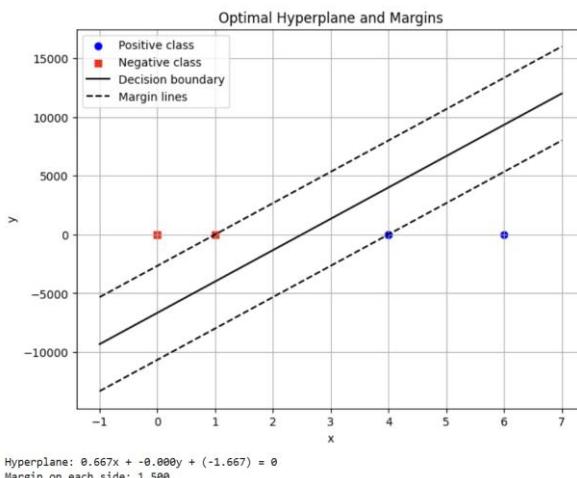
# Compute decision boundary and margins
xx = np.linspace(-1, 7, 500)
yy = -(w[0] * xx + b) / w[1]

# Margin offset: distance = 1/||w||
margin = 1 / np.linalg.norm(w)
yy_down = yy - np.sqrt(1 + (w[0] / w[1])**2) * margin
yy_up = yy + np.sqrt(1 + (w[0] / w[1])**2) * margin

# Plotting
plt.figure(figsize=(8, 6))
plt.scatter(X[y == 1, 0], X[y == 1, 1], c='blue', marker='o', label='Positive class')
plt.scatter(X[y == -1, 0], X[y == -1, 1], c='red', marker='s', label='Negative class')
plt.plot(xx, yy, 'k-', label='Decision boundary')
plt.plot(xx, yy_down, 'k--', label='Margin lines')
plt.plot(xx, yy_up, 'k--')
plt.xlabel('x')
plt.ylabel('y')
plt.legend()
plt.title('Optimal Hyperplane and Margins')
plt.grid(True)
plt.show()

# Print hyperplane equation
print(f"Hyperplane: {w[0]:.3f}x + {w[1]:.3f}y + ({b:.3f}) = 0")
print(f"Margin on each side: {margin:.3f}")

```



```
import pandas as pd
```

```
# Load both datasets
```

```

iris_df = pd.read_csv("/content/iris.csv")
# 1. IRIS DATASET - SVM with RBF and Linear Kernels
X_iris = iris_df.drop("species", axis=1)
y_iris = iris_df["species"]

# Encode labels
le_iris = LabelEncoder()
y_iris_encoded = le_iris.fit_transform(y_iris)

# Split dataset
X_train_iris, X_test_iris, y_train_iris, y_test_iris = train_test_split(X_iris, y_iris_encoded, test_size=0.2, random_state=42)

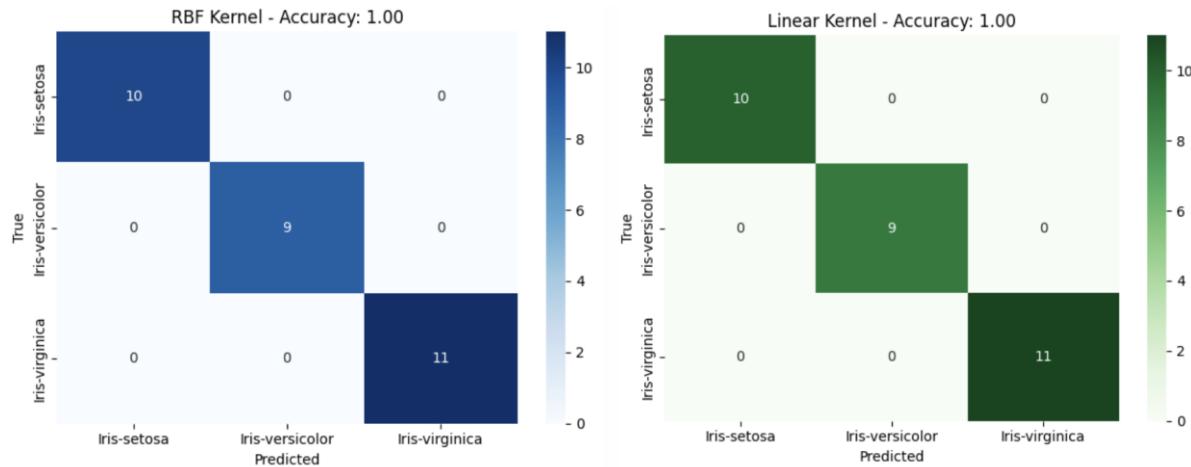
# Train models
svm_rbf = SVC(kernel='rbf')
svm_linear = SVC(kernel='linear')

svm_rbf.fit(X_train_iris, y_train_iris)
svm_linear.fit(X_train_iris, y_train_iris)

# Predictions
y_pred_rbf = svm_rbf.predict(X_test_iris)
y_pred_linear = svm_linear.predict(X_test_iris)

# Accuracy and Confusion Matrix
acc_rbf = accuracy_score(y_test_iris, y_pred_rbf)
acc_linear = accuracy_score(y_test_iris, y_pred_linear)
cm_rbf = confusion_matrix(y_test_iris, y_pred_rbf)
cm_linear = confusion_matrix(y_test_iris, y_pred_linear)

```



```

# Load dataset
letter_df = pd.read_csv("/content/letter-recognition.csv") # Update path if needed
letter_df['letter'] = LabelEncoder().fit_transform(letter_df['letter'])

# Split features and labels
X = letter_df.drop('letter', axis=1)
y = letter_df['letter']

# Train-test split
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)

# Standardize
scaler = StandardScaler()
X_train = scaler.fit_transform(X_train)
X_test = scaler.transform(X_test)

# Train SVM
svm = SVC(kernel='rbf', probability=True)
svm.fit(X_train, y_train)
y_pred = svm.predict(X_test)

```

```

y_prob = svm.predict_proba(X_test)

# Accuracy and Confusion Matrix
print("Accuracy:", accuracy_score(y_test, y_pred))
print("Confusion Matrix:\n", confusion_matrix(y_test, y_pred))

# ROC and AUC (one-vs-rest)
y_test_bin = label_binarize(y_test, classes=np.unique(y))
n_classes = y_test_bin.shape[1]

fpr = dict()
tpr = dict()
roc_auc = dict()

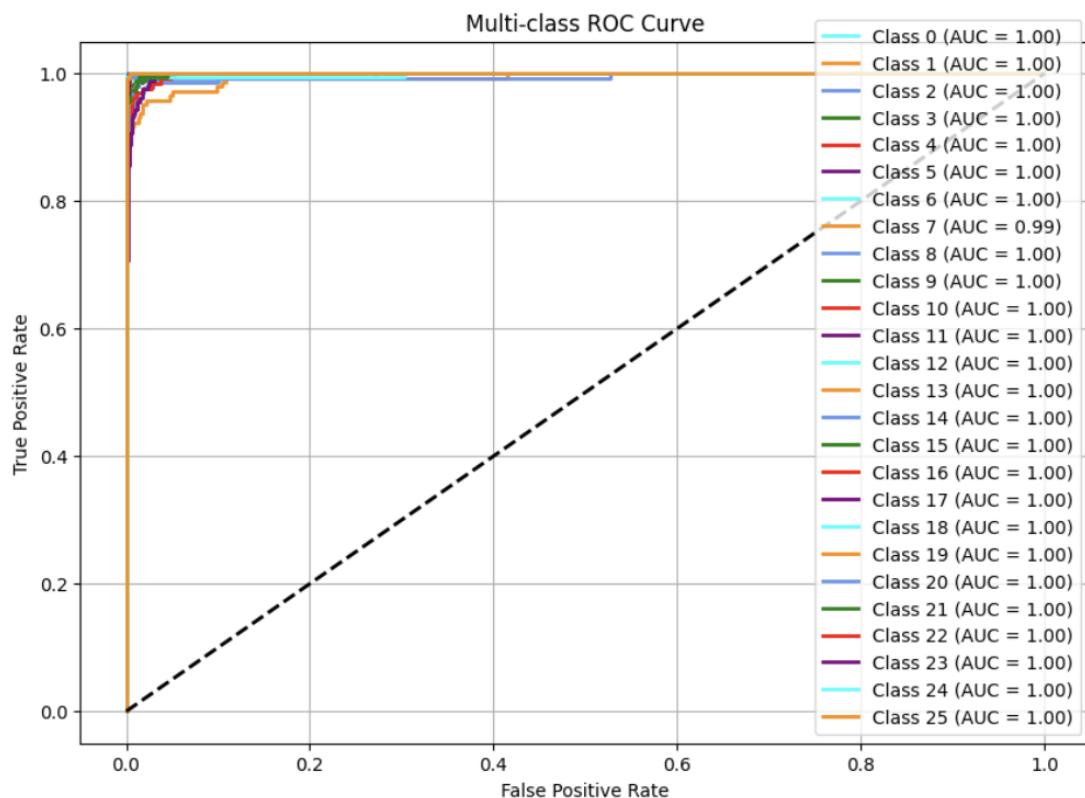
for i in range(n_classes):
    fpr[i], tpr[i], _ = roc_curve(y_test_bin[:, i], y_prob[:, i])
    roc_auc[i] = auc(fpr[i], tpr[i])

# Plot ROC Curve
plt.figure(figsize=(10, 7))
colors = cycle(['aqua', 'darkorange', 'cornflowerblue', 'green', 'red', 'purple'])

for i, color in zip(range(n_classes), colors):
    plt.plot(fpr[i], tpr[i], color=color, lw=2,
             label=f'Class {i} (AUC = {roc_auc[i]:0.2f})')

plt.plot([0, 1], [0, 1], 'k--', lw=2)
plt.xlabel("False Positive Rate")
plt.ylabel("True Positive Rate")
plt.title("Multi-class ROC Curve")
plt.legend(loc="lower right")
plt.grid()
plt.show()

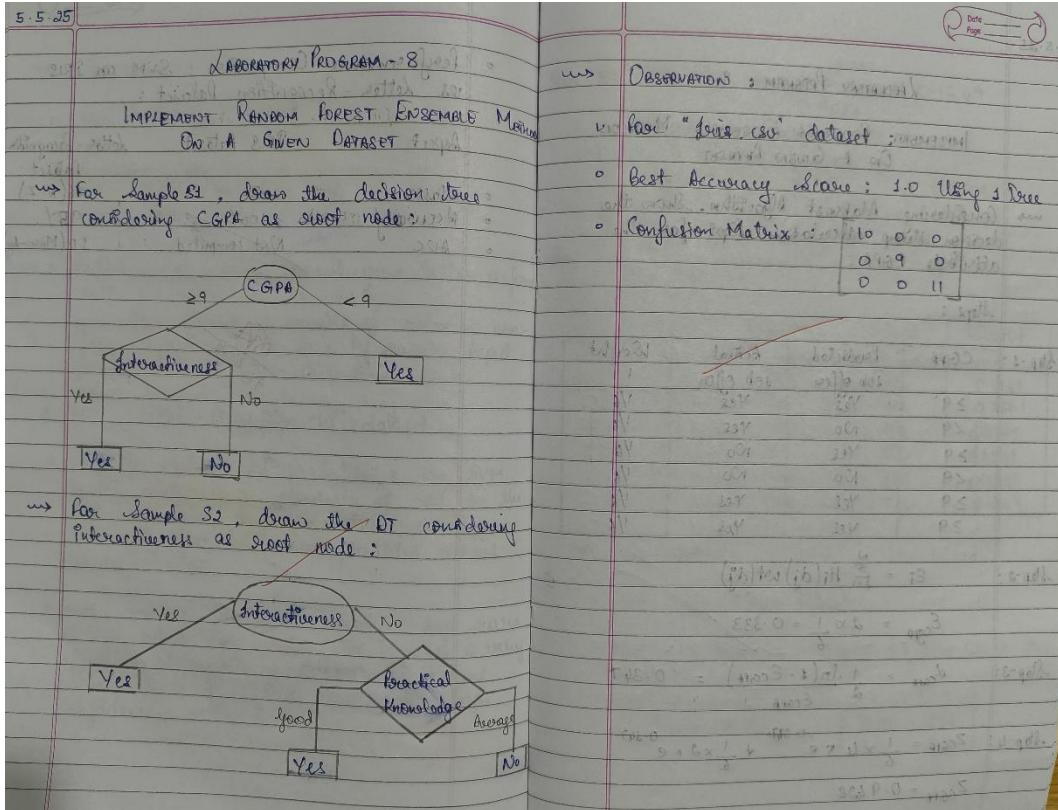
```



LABORATORY PROGRAM – 8

Implement Random forest ensemble method on a given dataset.

OBSERVATION BOOK



CODE WITH OUTPUT

```

import pandas as pd
from sklearn.model_selection import train_test_split
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import accuracy_score, confusion_matrix
import matplotlib.pyplot as plt

# Load the dataset
df = pd.read_csv("iris.csv") # Adjust filename if needed

# Prepare data
X = df.drop(columns=["species"]) # Assuming 'species' is the target column
y = df["species"]

# Split dataset
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)

# Default Random Forest with 10 trees
rf_default = RandomForestClassifier(n_estimators=10, random_state=42)
rf_default.fit(X_train, y_train)
y_pred_default = rf_default.predict(X_test)
acc_default = accuracy_score(y_test, y_pred_default)
conf_matrix_default = confusion_matrix(y_test, y_pred_default)

print(f"Default RF (10 trees) Accuracy: {acc_default}")
print("Confusion Matrix:\n", conf_matrix_default)

# Try different numbers of trees to find the best
  
```

```

best_acc = 0
best_n = 10
acc_list = []

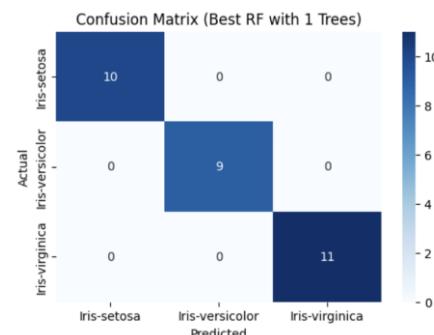
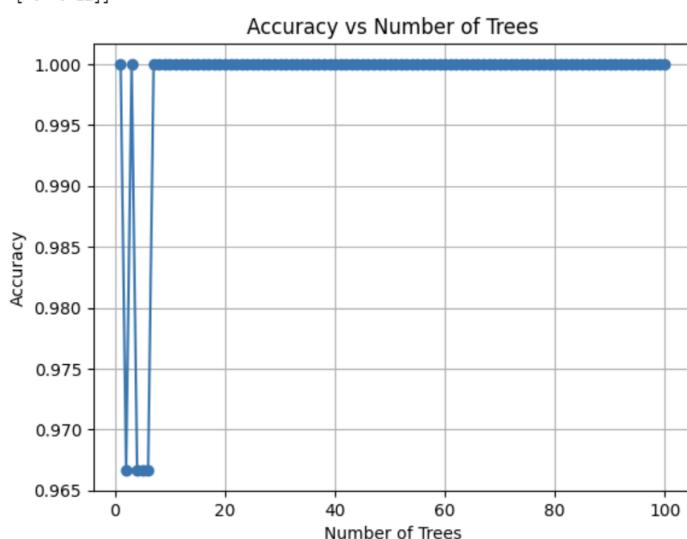
for n in range(1, 101):
    rf = RandomForestClassifier(n_estimators=n, random_state=42)
    rf.fit(X_train, y_train)
    y_pred = rf.predict(X_test)
    acc = accuracy_score(y_test, y_pred)
    acc_list.append((n, acc))
    if acc > best_acc:
        best_acc = acc
        best_n = n
        best_conf_matrix = confusion_matrix(y_test, y_pred)

print(f"\nBest Accuracy: {best_acc} using {best_n} trees")
print("Best Confusion Matrix:\n", best_conf_matrix)
# Plot accuracy vs number of trees
x_vals, y_vals = zip(*acc_list)
plt.plot(x_vals, y_vals, marker='o')
plt.title("Accuracy vs Number of Trees")
plt.xlabel("Number of Trees")
plt.ylabel("Accuracy")
plt.grid(True)
plt.show()

Default RF (10 trees) Accuracy: 1.0
Confusion Matrix:
[[10  0  0]
 [ 0  9  0]
 [ 0  0 11]]

Best Accuracy: 1.0 using 1 trees
Best Confusion Matrix:
[[10  0  0]
 [ 0  9  0]
 [ 0  0 11]]

```



LABORATORY PROGRAM – 9

Implement Boosting ensemble method on a given dataset.

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LABORATORY PROGRAM - 9 (INFORMATION)

**IMPLEMENT BOOSTING ENSEMBLE METHOD
ON A GIVEN DATASET**

using AdaBoost Algorithm. Show the decision stump calculation steps for the attribute CGPA.

Step :

Step-1:	CGPA	Predicted Job Offer	Actual Job Offer	Weight
	≥ 9	Yes	Yes	1/6
	< 9	No	Yes	1/6
	≥ 9	Yes	No	1/6
	< 9	No	No	1/6
	≥ 9	Yes	Yes	1/6
	≥ 9	Yes	Yes	1/6

Step-2: $E_i = \sum_{j=1}^N H_j(d_j) w_j(d_j)$

$E_{\text{GPA}} = 2 \times \frac{1}{6} = 0.333$

Step-3: $\alpha_{\text{base}} = \frac{1}{2} \ln(1 - E_{\text{GPA}}) = 0.347$

Step-4: $Z_{\text{CGPA}} = \frac{1}{6} \times 4 \times e^{-0.347} + \frac{1}{6} \times 2 \times e^{0.347}$

$Z_{\text{GPA}} = 0.9428$

Step-5: $w_j(d_j)_{\text{new}} = \frac{1}{6} \times e^{-0.347} = 0.247$
 0.9428

Final weight: $w_j(d_j)_{\text{final}} = \frac{1}{6} \times e^{0.347} = 0.2501$

for Incorrect instance: 0.9428

The Table:

CGPA	Predicted Job Offer	Actual Job Offer	Weight
≥ 9	Yes	Yes	0.1249
< 9	No	Yes	0.2501
≥ 9	Yes	No	0.1249
< 9	No	No	0.1249
≥ 9	Yes	Yes	0.1249
> 9	Yes	Yes	0.1249

Observations:

- for "income.csv" dataset over fit
- Best CV Accuracy: 0.8330 [5-fold CV on the training set].
- Test Accuracy: 0.8344.
- Confusion Matrix: $\begin{bmatrix} 7133 & 298 \\ 1320 & 1018 \end{bmatrix}$

CODE WITH OUTPUT

```

import pandas as pd
import matplotlib.pyplot as plt
from sklearn.model_selection import train_test_split, cross_val_score
from sklearn.preprocessing import OneHotEncoder, StandardScaler
from sklearn.compose import ColumnTransformer
from sklearn.pipeline import Pipeline
from sklearn.ensemble import AdaBoostClassifier
from sklearn.metrics import accuracy_score, confusion_matrix, ConfusionMatrixDisplay

# Load dataset
data = pd.read_csv('income.csv')

# Display basic info
print("First five rows:")
print(data.head())
print(f"\nDataset shape: {data.shape}")

# Define features and target
target_column = 'income_level'
y = data[target_column]
X = data.drop(columns=[target_column])

# Identify categorical vs numerical columns
categorical_cols = X.select_dtypes(include=['object', 'category']).columns.tolist()
numerical_cols = X.select_dtypes(include=['int64', 'float64']).columns.tolist()
print(f"\nNumerical columns: {numerical_cols}")
print(f"Categorical columns: {categorical_cols}")

```

```

# Preprocessor: scale numericals, one-hot encode categoricals
preprocessor = ColumnTransformer(
    transformers=[]
        ('num', StandardScaler(), numerical_cols),
        ('cat', OneHotEncoder(handle_unknown='ignore'), categorical_cols)
    )
)

# Initial AdaBoost model with 10 estimators
pipeline = Pipeline([
    ('preprocess', preprocessor),
    ('clf', AdaBoostClassifier(n_estimators=10, random_state=42))
])

# Split into train/test sets
X_train, X_test, y_train, y_test = train_test_split(
    X, y, test_size=0.2, random_state=42, stratify=y
)

# Train and evaluate initial model
pipeline.fit(X_train, y_train)
y_pred = pipeline.predict(X_test)
initial_acc = accuracy_score(y_test, y_pred)
print(f"Initial test accuracy (n_estimators=10): {initial_acc:.4f}")

# Hyperparameter tuning: find best n_estimators
tree_counts = list(range(10, 201, 10)) # 10,20,...,200
cv_scores = []
for n in tree_counts:
    model = Pipeline([
        ('preprocess', preprocessor),
        ('clf', AdaBoostClassifier(n_estimators=n, random_state=42))
    ])
    scores = cross_val_score(
        model, X_train, y_train, cv=5, scoring='accuracy', n_jobs=-1
    )
    mean_score = scores.mean()
    cv_scores.append(mean_score)
    print(f"n_estimators={n}: CV mean accuracy={mean_score:.4f}")

# Plot CV accuracy vs. number of estimators
plt.figure()
plt.plot(tree_counts, cv_scores, marker='o')
plt.title('AdaBoost CV Accuracy vs. n_estimators')
plt.xlabel('Number of Estimators')
plt.ylabel('CV Mean Accuracy')
plt.grid(True)
plt.tight_layout()
plt.show()

# Determine optimal number of trees
best_score = max(cv_scores)
best_n = tree_counts[cv_scores.index(best_score)]
print(f"\nBest CV accuracy={best_score:.4f} with n_estimators={best_n}")

# Retrain and evaluate best model
best_model = Pipeline([
    ('preprocess', preprocessor),
    ('clf', AdaBoostClassifier(n_estimators=best_n, random_state=42))
])
best_model.fit(X_train, y_train)
y_best = best_model.predict(X_test)
best_test_acc = accuracy_score(y_test, y_best)
print(f"Test accuracy with best n_estimators ({best_n}): {best_test_acc:.4f}")

# Plot comparison of initial vs. best test accuracy

```

```

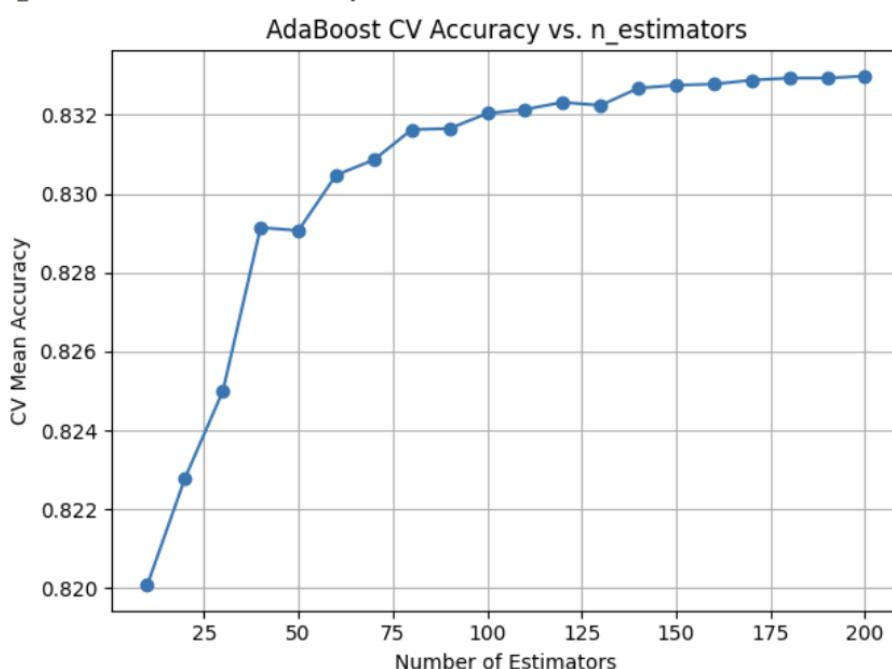
plt.figure()
plt.bar(['n=10', 'fn={best_n}'], [initial_acc, best_test_acc])
plt.title('Test Accuracy: Initial vs. Optimized')
plt.ylabel('Accuracy')
plt.ylim(0, 1)
plt.tight_layout()
plt.show()

# Plot confusion matrix for best model
cm = confusion_matrix(y_test, y_best)
labels = best_model.named_steps['clf'].classes_
disp = ConfusionMatrixDisplay(confusion_matrix=cm, display_labels=labels)
plt.figure()
disp.plot(cmap=plt.cm.Blues)
plt.title('Confusion Matrix for Best AdaBoost Model')
plt.tight_layout()
plt.show()

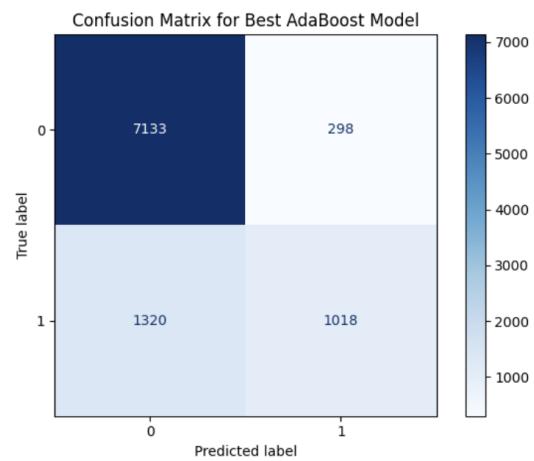
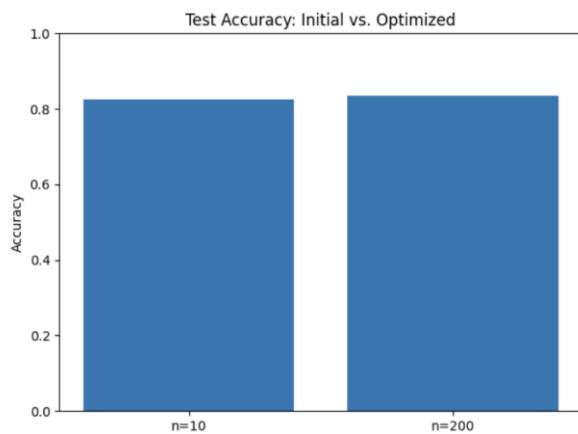
Dataset shape: (48842, 7)

Numerical columns: ['age', 'fnlwgt', 'education_num', 'capital_gain', 'capital_loss', 'hours_per_week']
Categorical columns: []
Initial test accuracy (n_estimators=10): 0.8257
n_estimators=10: CV mean accuracy=0.8201
n_estimators=20: CV mean accuracy=0.8228
n_estimators=30: CV mean accuracy=0.8250
n_estimators=40: CV mean accuracy=0.8291
n_estimators=50: CV mean accuracy=0.8291
n_estimators=60: CV mean accuracy=0.8305
n_estimators=70: CV mean accuracy=0.8309
n_estimators=80: CV mean accuracy=0.8316
n_estimators=90: CV mean accuracy=0.8316
n_estimators=100: CV mean accuracy=0.8320
n_estimators=110: CV mean accuracy=0.8321
n_estimators=120: CV mean accuracy=0.8323
n_estimators=130: CV mean accuracy=0.8322
n_estimators=140: CV mean accuracy=0.8327
n_estimators=150: CV mean accuracy=0.8327
n_estimators=160: CV mean accuracy=0.8328
n_estimators=170: CV mean accuracy=0.8329
n_estimators=180: CV mean accuracy=0.8329
n_estimators=190: CV mean accuracy=0.8329
n_estimators=200: CV mean accuracy=0.8330

```



Best CV accuracy=0.8330 with n_estimators=200
Test accuracy with best n_estimators (200): 0.8344



LABORATORY PROGRAM – 10

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

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Build K-Means Algorithm To CLUSTER A SET OF DATA STORED IN A .CSV FILE

→ Iteration 1: Record 1 to Cluster 1, Cluster 1: (1.83, 2.33) Assign to cluster
Record 2 to Cluster 1, Cluster 1: (4.12, 5.37) Assign to cluster
Record 3 to Cluster 2, Cluster 2: (3.61, 1.61) Assign to cluster
Record 4 to Cluster 2, Cluster 2: (5.64, 1.85) Assign to cluster
Record 5 to Cluster 2, Cluster 2: (3.15, 0.72) Assign to cluster
Record 6 to Cluster 2, Cluster 2: (3.78, 0.53) Assign to cluster
Record 7 to Cluster 2, Cluster 2: (2.74, 1.07) Assign to cluster

Therefore new cluster are:
Cluster 1: R₁, R₂ and Cluster 2: R₃, R₄, R₅, R₆, R₇

Their new centroid are:
 $C_1 = \frac{1.83 + 2.33}{2}, \frac{4.12 + 5.37}{2} = 2.08, 4.74$
 $C_2 = \frac{3.61 + 1.61 + 5.64 + 1.85 + 3.15 + 0.72 + 3.78}{7}, \frac{0.53 + 1.07}{7} = 3.9, 5.1$

Observations:
for "Iris.csv" dataset:

The elbow plot (Intertia vs k) shows a sharp "elbow" at k=3, indicating that three clusters is the optimal choice for the petal-length/width of Iris dataset.

CODE WITH OUTPUT

```

import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
from sklearn.datasets import load_iris
from sklearn.preprocessing import StandardScaler
from sklearn.cluster import KMeans
from sklearn.metrics import confusion_matrix, ConfusionMatrixDisplay

def load_data(csv_path='iris.csv'):
    """
    Try loading from csv_path; if not found, load via sklearn.
    Expects columns: sepal_length, sepal_width, petal_length, petal_width, species.
    Returns DataFrame with a 'species' column.
    """
    try:
        df = pd.read_csv(csv_path)
        # Fixed typo here: use c.strip().replace, not ace()
        df.columns = [c.strip().replace(' ', '_') for c in df.columns]
    except FileNotFoundError:
        iris = load_iris()
        df = pd.DataFrame(
            data=np.c_[iris['data'], iris['target']],
            columns=iris['feature_names'] + ['target']
        )
        df.columns = [c.strip().replace(' (cm)', '').replace(' ', '_')
                     for c in df.columns]

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df['species'] = df['target'].map(lambda x: iris['target_names'][int(x)])
return df

def preprocess(df):
    """
    Select only petal_length & petal_width, then standard-scale.
    Returns scaled numpy array.
    """
    X = df[['petal_length', 'petal_width']].values
    scaler = StandardScaler()
    X_scaled = scaler.fit_transform(X)
    return X_scaled, scaler

def plot_elbow(X_scaled, max_k=10):
    """
    Compute KMeans inertia for k=1..max_k and plot the elbow curve.
    Returns list of inertias.
    """
    inertias = []
    ks = range(1, max_k + 1)
    for k in ks:
        km = KMeans(n_clusters=k, random_state=42)
        km.fit(X_scaled)
        inertias.append(km.inertia_)
    plt.figure(figsize=(6, 4))
    plt.plot(ks, inertias, 'o-', linewidth=2)
    plt.xlabel('Number of clusters (k)')
    plt.ylabel('Inertia')
    plt.title('Elbow Method for Optimal k')
    plt.xticks(ks)
    plt.grid(True, linestyle='--', alpha=0.5)
    plt.tight_layout()
    plt.show()
    return inertias

def run_kmeans(X_scaled, k):
    """
    Fit KMeans with k clusters, return labels and fitted model.
    """
    km = KMeans(n_clusters=k, random_state=42)
    labels = km.fit_predict(X_scaled)
    return km, labels

def plot_confusion(df, labels, k):
    """
    Builds and displays a confusion matrix comparing true species vs. cluster.
    """
    species_names = df['species'].unique()
    species_to_num = {name: idx for idx, name in enumerate(species_names)}
    true_nums = df['species'].map(species_to_num)

    cm = confusion_matrix(true_nums, labels)
    disp = ConfusionMatrixDisplay(
        confusion_matrix=cm,
        display_labels=[f"Cluster {i}" for i in range(k)])
    fig, ax = plt.subplots(figsize=(6, 6))
    disp.plot(ax=ax, cmap='Blues', colorbar=True)
    ax.set_xlabel('Predicted Cluster')
    ax.set_ylabel('True Species')
    plt.title('K-Means Clustering Confusion Matrix')
    plt.tight_layout()
    plt.show()

    cm_df = pd.DataFrame(
        cm,
        index=[f"True: {name}" for name in species_names],

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        columns=[f"Cluster {i}" for i in range(k)]
    )
print("\nConfusion Matrix (counts):")
print(cm_df)

def main():
    # 1) Load data
    df = load_data('iris.csv')
    if 'species' not in df.columns:
        print("Error: 'species' column not found.")
        return

    # 2) Preprocess
    X_scaled, scaler = preprocess(df)

    # 3) Elbow plot to decide k
    print("Generating elbow plot to find optimal k...")
    inertias = plot_elbow(X_scaled, max_k=10)

    # 4) From the elbow you'll typically see a bend at k=3
    optimal_k = 3
    print(f"Choosing k = {optimal_k} (you can adjust this based on the plot.)")

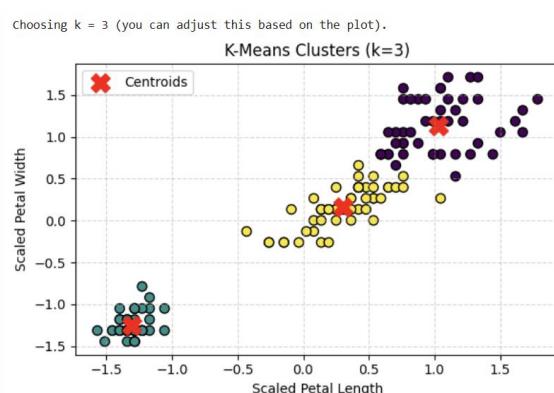
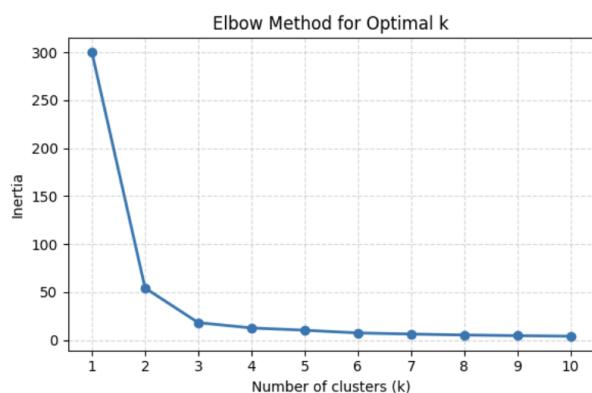
    # 5) Run K-Means and assign clusters
    km_model, labels = run_kmeans(X_scaled, optimal_k)
    df['cluster'] = labels

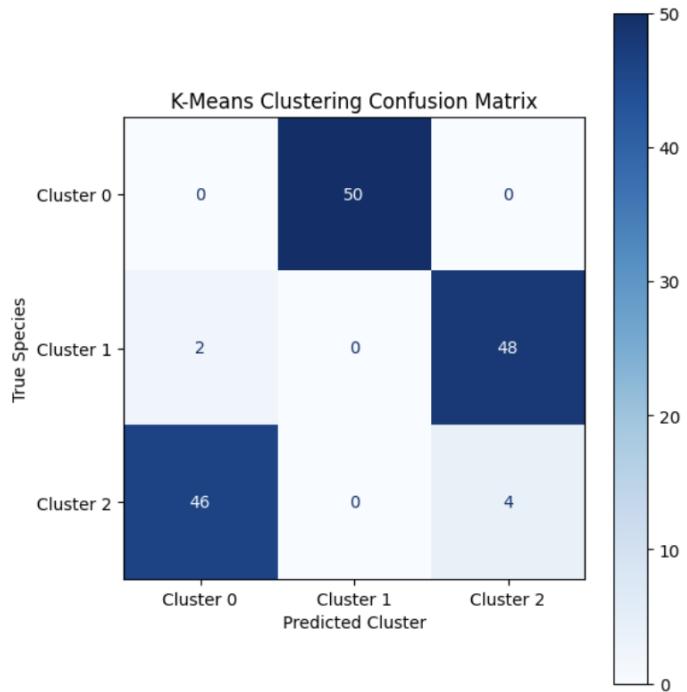
    # 6) Visualize clusters in feature space
    plt.figure(figsize=(6, 4))
    plt.scatter(
        X_scaled[:, 0], X_scaled[:, 1],
        c=labels, cmap='viridis', edgecolor='k', s=50
    )
    centroids = km_model.cluster_centers_
    plt.scatter(
        centroids[:, 0], centroids[:, 1],
        marker='X', c='red', s=200, label='Centroids'
    )
    plt.xlabel('Scaled Petal Length')
    plt.ylabel('Scaled Petal Width')
    plt.title(f'K-Means Clusters (k={optimal_k})')
    plt.legend()
    plt.grid(True, linestyle='--', alpha=0.5)
    plt.tight_layout()
    plt.show()

    # 7) Confusion matrix vs. true species
    plot_confusion(df, labels, optimal_k)

if __name__ == "__main__":
    main()

```





LABORATORY PROGRAM – 11

Implement Dimensionality reduction using Principle Component Analysis (PCA) method.

OBSERVATION BOOK

5.5.25	LABORATORY PROGRAM - 11	Z ₁ (-4, -2.5) = 0.5574(-4) + (-0.8303)(-2.5) = -4.30535
5.5.25	IMPLEMENT DIMENSIONALITY REDUCTION USING PRINCIPLE COMPONENT ANALYSIS (PCA) METHOD	Z ₂ (0, -4.5) = 0.5574(0) + (-0.8303)(-4.5) = 3.73635
5.5.25	→ Reduce the dimension from 2 to 1 using the PCA. Compute for principal component.	Z ₃ (5, -3.5) = 0.5574(5) + (-0.8303)(-3.5) = 5.69305
5.5.25	→ Data Matrix :	Z ₄ (-1, 5.5) = 0.5574(-1) + (-0.8303)(5.5) = -5.12405
5.5.25	X ₁ 4 8 18 7 9 X ₂ 11 4 5 14	Z = [-4.305] [3.736] [5.693] [-5.124]
5.5.25	Mean of X ₁ = $\frac{4+18+18+7}{4} = 12.25$ Mean of X ₂ = $\frac{11+4+5+14}{4} = 8.5$ $\bar{X}_{\text{combined}} = \bar{X} - \text{Mean} = \begin{bmatrix} -4 & 0 & 5 & -1 \\ 2.5 & -4.5 & -3.5 & 5.5 \end{bmatrix}$	ms OBSERVATION
5.5.25	Since λ_1 is larger, e ₁ is the first Principal Component	1. For "heart.csv" dataset : • Accuracy Before PCA : Logistic Regression : 0.9016 SVM : 0.8525 Random Forest : 0.8361
5.5.25	e ₁ = [0.5574 to -0.8303] Let Z _i = e ₁ · x _i	• Accuracy After PCA : Logistic Regression : 0.8689 SVM : 0.8689 Random Forest : 0.8852

CODE WITH OUTPUT

```
import pandas as pd  
  
df = pd.read_csv("heart.csv")  
  
# Step 3: Split Features and Target  
X = df.drop("target", axis=1)  
y = df["target"]  
  
# Step 4: Preprocessing  
from sklearn.model_selection import train_test_split  
from sklearn.preprocessing import StandardScaler, OneHotEncoder  
from sklearn.compose import ColumnTransformer  
from sklearn.pipeline import Pipeline  
  
categorical_features = ["cp", "thal", "slope"]  
numerical_features = [col for col in X.columns if col not in categorical_features]  
  
preprocessor = ColumnTransformer(transformers=[  
    ("num", StandardScaler(), numerical_features),  
    ("cat", OneHotEncoder(), categorical_features)  
])  
  
# Step 5: Train/Test Split  
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
```

```

# Step 6: Models
from sklearn.linear_model import LogisticRegression
from sklearn.svm import SVC
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import accuracy_score

models = {
    "Logistic Regression": LogisticRegression(max_iter=1000),
    "SVM": SVC(),
    "Random Forest": RandomForestClassifier()
}

# Step 7: Train and Evaluate Models (Before PCA)
print("Accuracy Before PCA:")
results = {}
for name, model in models.items():
    pipeline = Pipeline(steps=[
        ("preprocessor", preprocessor),
        ("classifier", model)
    ])
    pipeline.fit(X_train, y_train)
    y_pred = pipeline.predict(X_test)
    acc = accuracy_score(y_test, y_pred)
    results[name] = acc
    print(f"{name}: {acc:.4f}")

from sklearn.decomposition import PCA

print("\nAccuracy After PCA (n_components=5):")
pca_results = {}

for name, model in models.items():
    pipeline_pca = Pipeline(steps=[
        ("preprocessor", preprocessor),
        ("pca", PCA(n_components=5)),
        ("classifier", model)
    ])
    pipeline_pca.fit(X_train, y_train)
    y_pred_pca = pipeline_pca.predict(X_test)
    acc_pca = accuracy_score(y_test, y_pred_pca)
    pca_results[name] = acc_pca
    print(f"{name}: {acc_pca:.4f}")

```

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→ └─ Accuracy Before PCA:
  Logistic Regression: 0.9016
  SVM: 0.8525
  Random Forest: 0.8361

  └─ Accuracy After PCA (n_components=5):
  Logistic Regression: 0.8689
  SVM: 0.8689
  Random Forest: 0.8852

```