

MACHINE LEARNING

A PRACTICAL REPORT ON
MACHINE LEARNING

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Seat No :

UNDER THE GUIDANCE OF
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Department of Information Technology

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Certificate

This is to certify that on Machine Learning Practical's
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College Stamp

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Practical No: - 1

Aim: Data Pre-processing and Exploration.

Writeups:

[illegible]

1A) Aim: Load a CSV dataset. Handle missing values, inconsistent formatting, and outliers.

Code:

```
import pandas as pd
import numpy as np
import requests
from io import StringIO
from scipy import stats
url = 'https://archive.ics.uci.edu/ml/machine-learning-
databases/iris/iris.data'
columns = ['sepal_length', 'sepal_width', 'petal_length', 'petal_width',
'species']
response = requests.get(url)
csv_data = StringIO(response.text)
df = pd.read_csv(csv_data, header=None, names=columns)
print("Before Data Cleaning:")
print(df.info())
print(df.head())
df_imputed = df.copy()
numeric_cols = df_imputed.select_dtypes(include=np.number).columns
df_imputed[numeric_cols] =
df_imputed[numeric_cols].fillna(df_imputed[numeric_cols].mean())
categorical_cols = df_imputed.select_dtypes(include='object').columns
for col in categorical_cols:
    df_imputed[col].fillna(df_imputed[col].mode()[0], inplace=True)
df_imputed.columns = df_imputed.columns.str.strip().str.lower().str.replace('
', '_')
z_scores = np.abs(stats.zscore(df_imputed.select_dtypes(include=np.number)))
outliers = (z_scores > 3)
df_cleaned = df_imputed[~np.any(outliers, axis=1)]
print("\nAfter Data Cleaning:")
print(df_cleaned.info())
print(df_cleaned.head())
print("\nSummary Statistics Before Cleaning:")
print(df.describe())
print("\nSummary Statistics After Cleaning:")
print(df_cleaned.describe())
```

Output:

```
PS C:\Users\Shivang Singh> python -u "c:\Users\Shivang Singh\prac1a.py"
Before Data Cleaning:
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 150 entries, 0 to 149
Data columns (total 5 columns):
#   Column          Non-Null Count  Dtype
---  ---
0   sepal_length    150 non-null   float64
1   sepal_width     150 non-null   float64
2   petal_length    150 non-null   float64
3   petal_width     150 non-null   float64
4   species         150 non-null   object
dtypes: float64(4), object(1)
memory usage: 6.0+ KB
None
```

	sepal_length	sepal_width	petal_length	petal_width	species
0	5.1	3.5	1.4	0.2	Iris-setosa
1	4.9	3.0	1.4	0.2	Iris-setosa
2	4.7	3.2	1.3	0.2	Iris-setosa
3	4.6	3.1	1.5	0.2	Iris-setosa
4	5.0	3.6	1.4	0.2	Iris-setosa

```
After Data Cleaning:
<class 'pandas.core.frame.DataFrame'>
Index: 149 entries, 0 to 149
Data columns (total 5 columns):
#   Column          Non-Null Count  Dtype
---  ---
0   sepal_length    149 non-null   float64
1   sepal_width     149 non-null   float64
2   petal_length    149 non-null   float64
3   petal_width     149 non-null   float64
4   species         149 non-null   object
dtypes: float64(4), object(1)
memory usage: 7.0+ KB
None
```

	sepal_length	sepal_width	petal_length	petal_width	species
0	5.1	3.5	1.4	0.2	Iris-setosa
1	4.9	3.0	1.4	0.2	Iris-setosa
2	4.7	3.2	1.3	0.2	Iris-setosa
3	4.6	3.1	1.5	0.2	Iris-setosa
4	5.0	3.6	1.4	0.2	Iris-setosa

Summary Statistics Before Cleaning:

	sepal_length	sepal_width	petal_length	petal_width
count	150.000000	150.000000	150.000000	150.000000
mean	5.843333	3.054000	3.758667	1.198667
std	0.828066	0.433594	1.764420	0.763161
min	4.300000	2.000000	1.000000	0.100000
25%	5.100000	2.800000	1.600000	0.300000
50%	5.800000	3.000000	4.350000	1.300000
75%	6.400000	3.300000	5.100000	1.800000
max	7.900000	4.400000	6.900000	2.500000

Summary Statistics After Cleaning:

	sepal_length	sepal_width	petal_length	petal_width
count	149.000000	149.000000	149.000000	149.000000
mean	5.844295	3.044966	3.773826	1.204027
std	0.830775	0.420655	1.760543	0.762896
min	4.300000	2.000000	1.000000	0.100000
25%	5.100000	2.800000	1.600000	0.300000
50%	5.800000	3.000000	4.400000	1.300000
75%	6.400000	3.300000	5.100000	1.800000
max	7.900000	4.200000	6.900000	2.500000

1B) Aim: Load a dataset, calculate descriptive summary statistics, create visualizations using different graphs, and identify potential features and target variables Note: Explore Univariate and Bivariate graphs (Matplotlib) and Seaborn for visualization.

```
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns

def load_dataset(file_url):
    try:
        data = pd.read_csv(file_url)
        print("Dataset loaded successfully!")
        return data
    except Exception as e:
        print(f"Error loading dataset: {e}")
        return None

def descriptive_statistics(data):
    print("\nDescriptive Statistics:")
    print(data.describe(include='all'))
    print("\nMissing Values in Each Column:")
    print(data.isnull().sum())

def create_visualizations(data):
    sns.set(style="whitegrid")

    # Filter only numeric columns for plotting
    numeric_data = data.select_dtypes(include='number')

    # Histogram
    numeric_data.hist(bins=30, edgecolor='black', figsize=(12, 10))
    plt.suptitle('Histogram of Numeric Features')
    plt.show()

    # Box Plot
    plt.figure(figsize=(12, 6))
    sns.boxplot(data=numeric_data)
    plt.title('Box Plot of Numeric Features')
    plt.xticks(rotation=45)
    plt.show()

    # Correlation Heatmap
    plt.figure(figsize=(10, 8))
    corr = numeric_data.corr()
    sns.heatmap(corr, annot=True, cmap='coolwarm', linewidths=0.5)
    plt.title('Correlation Heatmap of Numeric Features')
```

```
plt.show()

# Pairplot (use only selected numeric columns to avoid long processing or
errors)
selected_cols = ['Age', 'Fare', 'Pclass'] # Safe numeric columns from
Titanic dataset
sns.pairplot(data[selected_cols].dropna()) # Drop rows with NaN
plt.suptitle('🌀 Pairplot of Selected Numeric Features', y=1.02)
plt.show()

def main():
    file_url =
'https://raw.githubusercontent.com/datasciencedojo/datasets/master/titanic.csv'

    data = load_dataset(file_url)

    if data is not None:
        descriptive_statistics(data)
        create_visualizations(data)

if __name__ == "__main__":
    main()
```

Output:

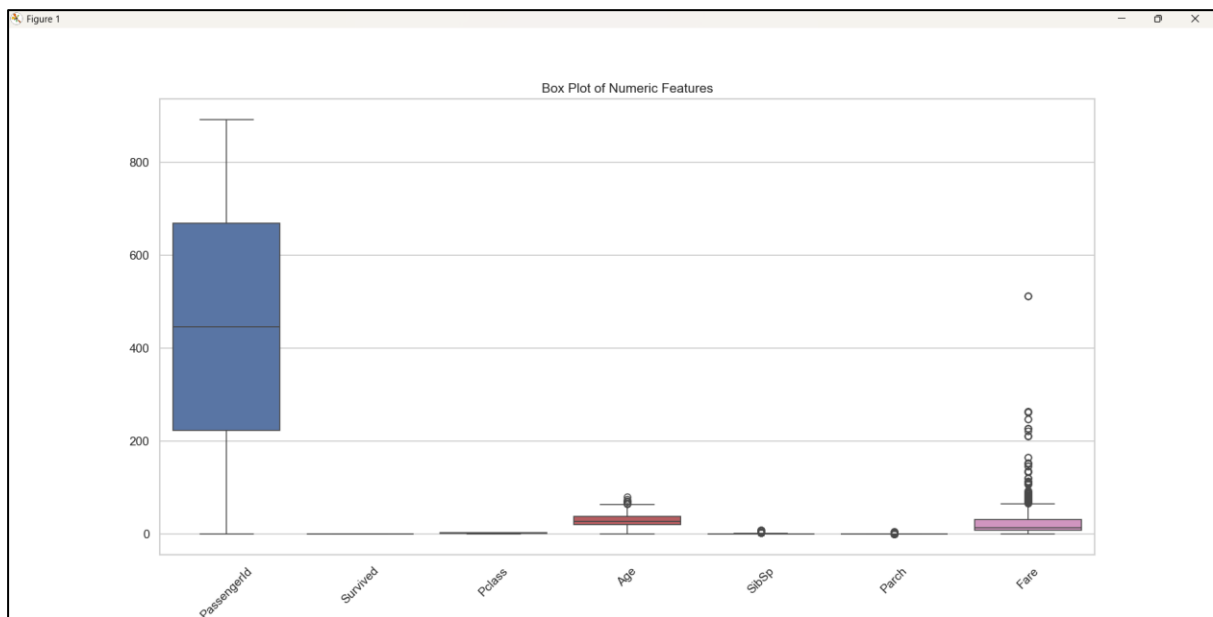
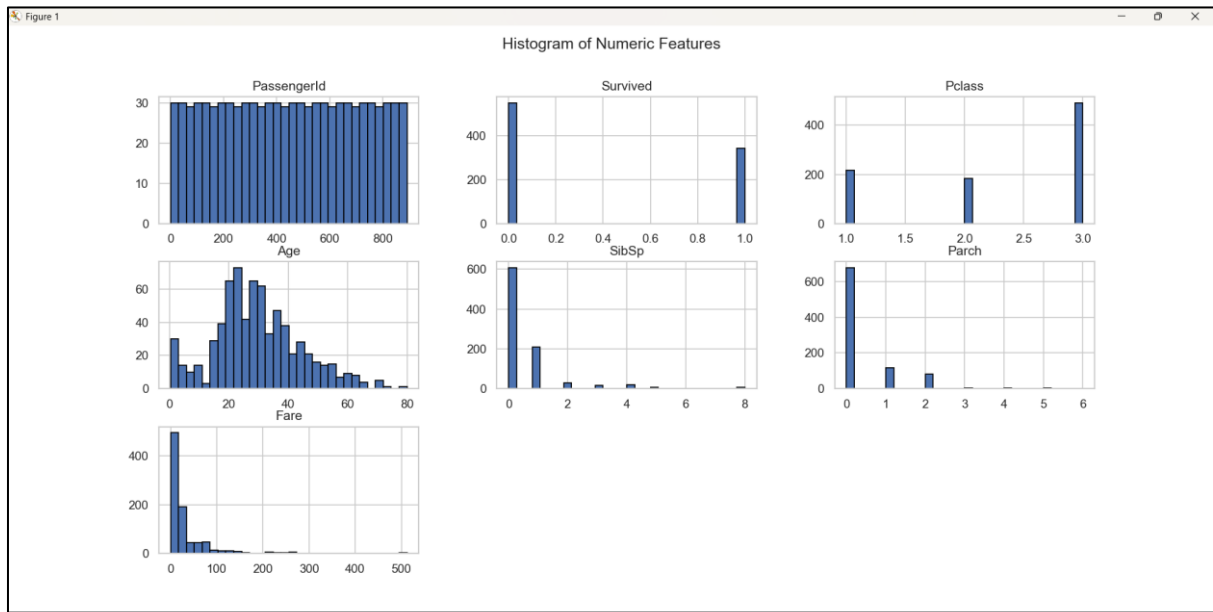
Dataset loaded successfully!

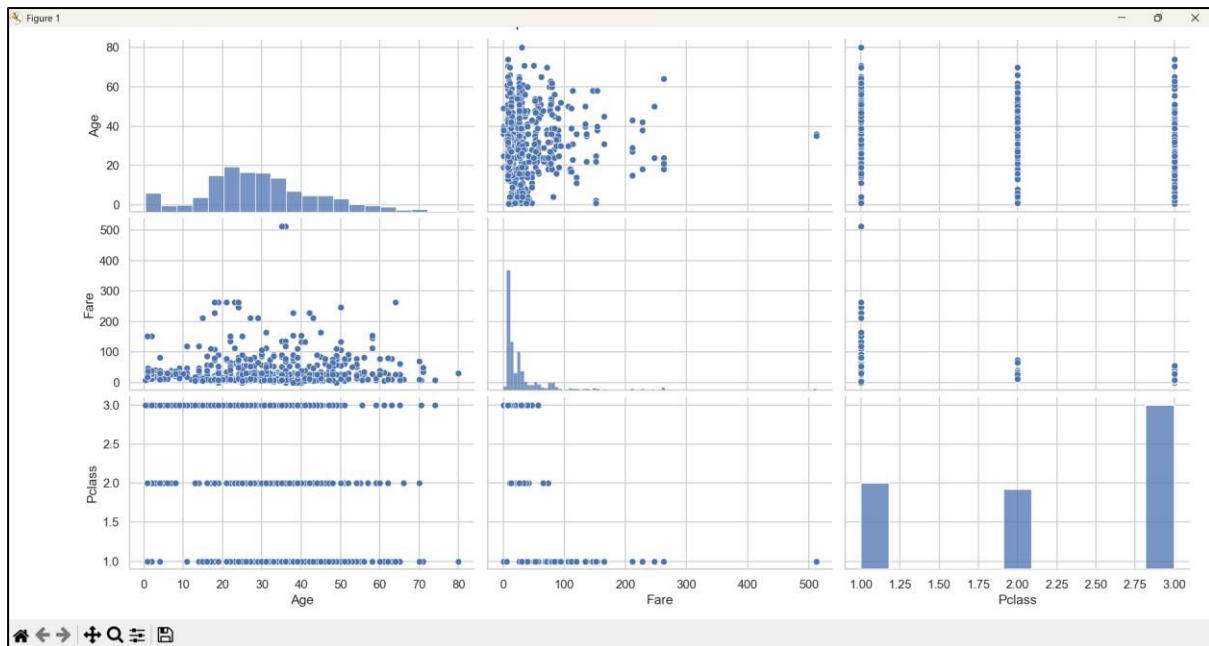
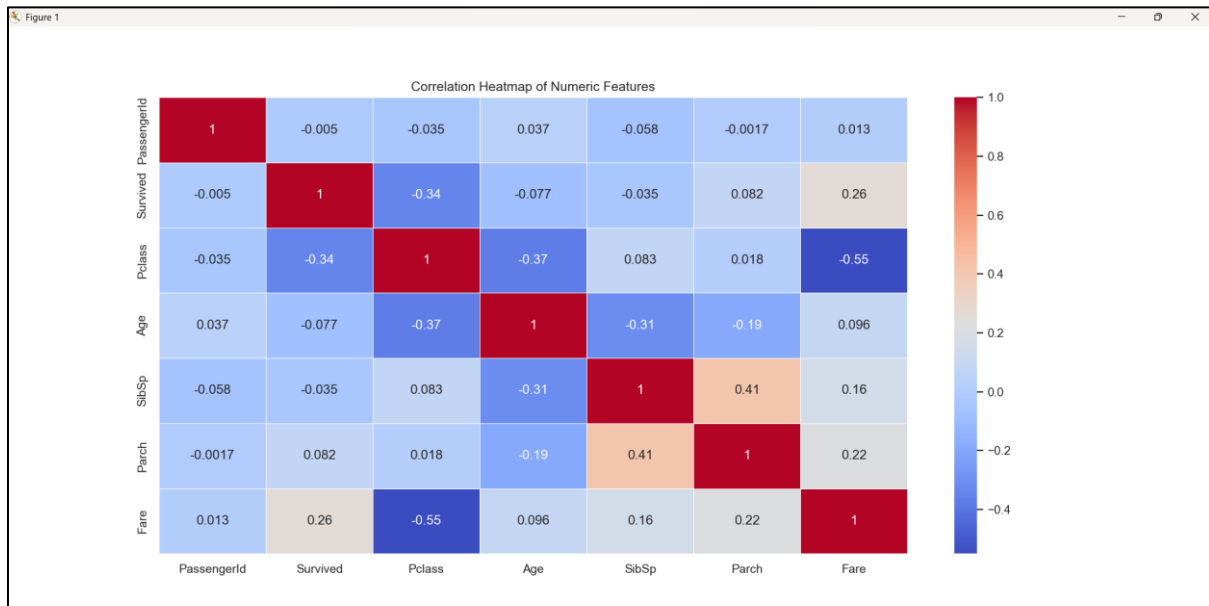
Descriptive Statistics:

	PassengerId	Survived	Pclass	Name	Sex	Age	SibSp	Parch	Ticket	Fare	Cabin	Embarked
count	891.000000	891.000000	891.000000	891	891	714.000000	891.000000	891.000000	891	891.000000	204	889
unique	NaN	NaN	NaN	891	2	NaN	NaN	NaN	681	NaN	147	3
top	NaN	NaN	NaN	Dooley, Mr. Patrick	male	NaN	NaN	NaN	347082	NaN	G6	S
freq	NaN	NaN	NaN	1	577	NaN	NaN	NaN	7	NaN	4	644
mean	446.000000	0.383838	2.308642	NaN	NaN	29.699118	0.523088	0.381594	NaN	32.204208	NaN	NaN
std	257.353842	0.486592	0.836071	NaN	NaN	14.526497	1.102743	0.806057	NaN	49.693429	NaN	NaN
min	1.000000	0.000000	1.000000	NaN	NaN	0.420000	0.000000	0.000000	NaN	0.000000	NaN	NaN
25%	223.500000	0.000000	2.000000	NaN	NaN	20.125000	0.000000	0.000000	NaN	7.910400	NaN	NaN
50%	446.000000	0.000000	3.000000	NaN	NaN	28.000000	0.000000	0.000000	NaN	14.454200	NaN	NaN
75%	668.500000	1.000000	3.000000	NaN	NaN	38.000000	1.000000	0.000000	NaN	31.000000	NaN	NaN
max	891.000000	1.000000	3.000000	NaN	NaN	80.000000	8.000000	6.000000	NaN	512.329200	NaN	NaN

Missing Values in Each Column:

```
PassengerId      0
Survived          0
Pclass           0
Name             0
Sex              0
Age             177
SibSp            0
Parch            0
Ticket           0
Fare             0
Cabin           687
Embarked         2
dtype: int64
```





1C) Aim: Create or Explore datasets to use all pre-processing routines like label encoding, scaling, and binarization.

Source Code

```
import pandas as pd
from sklearn.preprocessing import LabelEncoder, MinMaxScaler, Binarizer

def load_dataset(url):
    try:
        return pd.read_csv(url)
    except Exception as e:
        print(f"Error loading dataset: {e}")
        return None

def preprocess_data(data):
    numeric_cols = data.select_dtypes(include=['number']).columns
    data[numeric_cols] = data[numeric_cols].fillna(data[numeric_cols].mean())
    for col in data.select_dtypes(include='object').columns:
        data[col] = LabelEncoder().fit_transform(data[col].astype(str))
    scaler = MinMaxScaler()
    data[numeric_cols] = scaler.fit_transform(data[numeric_cols])
    binarizer = Binarizer(threshold=0.5)
    data[numeric_cols] = binarizer.fit_transform(data[numeric_cols])
    return data

def main():
    url = 'https://raw.githubusercontent.com/datasciencedojo/datasets/master/titanic.csv'

    data = load_dataset(url)
    if data is not None:
        print("Original Data:")
        print(data.head())
        data = preprocess_data(data)
        print("\nPreprocessed Data:")
        print(data.head())

if __name__ == "__main__":
    main()
```

Output:

```
Original Data:
 PassengerId  Survived  Pclass
0            1         0       3
1            2         1       1
2            3         1       3
3            4         1       1
4            5         0       3
      Name      Sex  Age  SibSp  Parch    Ticket   Fare Cabin Embarked
0  Braund, Mr. Owen Harris   male  22.0      1      0      A/5 21171   7.2500   NaN      S
1  Cumings, Mrs. John Bradley (Florence Briggs Th... female  38.0      1      0      PC 17599  71.2833   C85      C
2     Heikkinen, Miss. Laina   female  26.0      0      0  STON/O2. 3101282   7.9250   NaN      S
3  Futrelle, Mrs. Jacques Heath (Lily May Peel)   female  35.0      1      0     113803  53.1000  C123      S
4    Allen, Mr. William Henry   male  35.0      0      0     373450   8.0500   NaN      S

Preprocessed Data:
 PassengerId  Survived  Pclass  Name      Sex  Age  SibSp  Parch  Ticket   Fare  Cabin Embarked
0           0.0       0.0     1.0  108      1  0.0   0.0   0.0     523   0.0   147      2
1           0.0       1.0     0.0  190      0  0.0   0.0   0.0     596   0.0   81      0
2           0.0       1.0     1.0  353      0  0.0   0.0   0.0     669   0.0  147      2
3           0.0       1.0     0.0  272      0  0.0   0.0   0.0      49   0.0   55      2
4           0.0       0.0     1.0   15      1  0.0   0.0   0.0     472   0.0  147      2
```

Practical No: - 2

Aim: Hypothesis Testing.

Writeups:

[illegible]

Aim: Implement and demonstrate the FIND-S algorithm for finding the most specific hypothesis based on a given set of training data samples. Read the training data from a CSV file and generate the final specific hypothesis.

Code:

```
import pandas as pd

def find_s_algorithm(data, target_column):
    specific_hypothesis = ['0'] * (len(data.columns) - 1)
    for _, row in data.iterrows():
        if row[target_column] == "Yes":
            for i in range(len(specific_hypothesis)):
                if specific_hypothesis[i] == '0':
                    specific_hypothesis[i] = row.iloc[i]
                elif specific_hypothesis[i] != row.iloc[i]:
                    specific_hypothesis[i] = '?'
    return specific_hypothesis

def main():
    dataset = {
        'Sky': ['Sunny', 'Sunny', 'Rainy', 'Sunny', 'Sunny'],
        'Temp': ['Warm', 'Warm', 'Cold', 'Warm', 'Warm'],
        'Humidity': ['Normal', 'High', 'High', 'High', 'Normal'],
        'Wind': ['Strong', 'Strong', 'Strong', 'Strong', 'Strong'],
        'Water': ['Warm', 'Warm', 'Cold', 'Warm', 'Warm'],
        'Forecast': ['Same', 'Same', 'Change', 'Same', 'Same'],
        'EnjoySport': ['Yes', 'Yes', 'No', 'Yes', 'Yes']
    }
    df = pd.DataFrame(dataset)
    df.to_csv("training_data.csv", index=False)
    print("Dataset saved to training_data.csv")

    data = pd.read_csv("training_data.csv")
    print("\nTraining Data:")
    print(data)

    specific_hypothesis = find_s_algorithm(data, 'EnjoySport')
    print("\nFinal Specific Hypothesis:")
    print(specific_hypothesis)

if __name__ == "__main__":
    main()
```

Output:

```
PROBLEMS  OUTPUT  DEBUG CONSOLE  TERMINAL  PORTS

PS C:\Users\Shivang Singh> python -u "C:\Users\SHIVAN~1\AppData\Local\Temp\tempCodeRunnerFile.python"
Dataset saved to training_data.csv

Training Data:
   Sky  Temp Humidity  Wind Water Forecast EnjoySport
0  Sunny  Warm   Normal  Strong  Warm     Same      Yes
1  Sunny  Warm    High  Strong  Warm     Same      Yes
2  Rainy  Cold    High  Strong  Cold    Change     No
3  Sunny  Warm    High  Strong  Warm     Same      Yes
4  Sunny  Warm   Normal  Strong  Warm     Same      Yes

Final Specific Hypothesis:
['Sunny', 'Warm', '?', 'Strong', 'Warm', 'Same']
```


Practical No: - 3

Aim: Linear Models

Writeups:

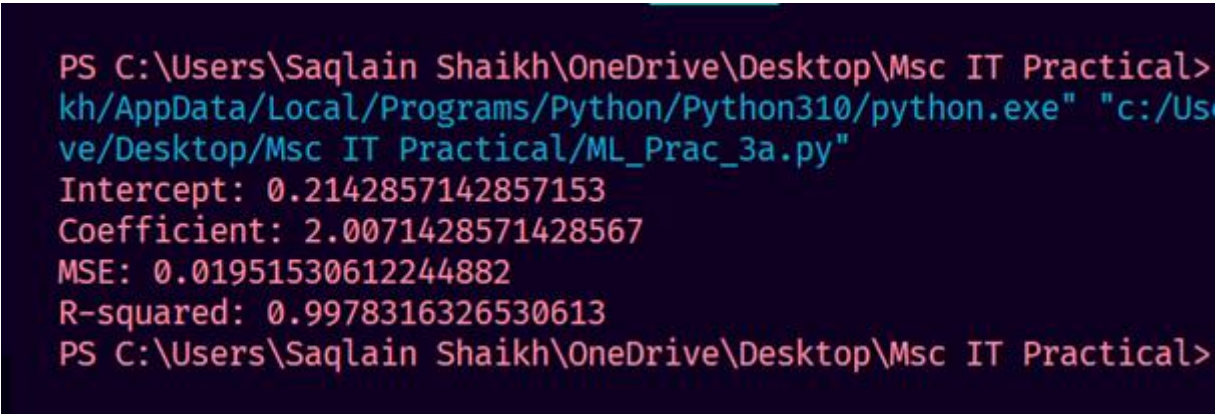
[illegible]

Practical 3A:**Code:**

```
import pandas as pd
from sklearn.model_selection import train_test_split
from sklearn.linear_model import LinearRegression
from sklearn.metrics import mean_squared_error, r2_score
data = {'X': [1, 2, 3, 4, 5], 'Y': [2.2, 4.1, 6.3, 8.2, 10.1]}
df = pd.DataFrame(data)
X = df[['X']]
Y = df[['Y']]
X_train, X_test, Y_train, Y_test = train_test_split(X, Y,
test_size=0.4,
random_state=42)
model = LinearRegression()
model.fit(X_train, Y_train)
intercept = model.intercept_
coefficient = model.coef_[0]
Y_pred = model.predict(X_test)
mse = mean_squared_error(Y_test, Y_pred)

if len(Y_test) > 1:
    r_squared = r2_score(Y_test, Y_pred)
else:
    r_squared = float('nan') # Set R-squared to NaN if only one
test sample

print(f"Intercept: {intercept}")
print(f"Coefficient: {coefficient}")
print(f"MSE: {mse}")
print(f"R-squared: {r_squared}")
```

OUTPUT:

```
PS C:\Users\Saqlain Shaikh\OneDrive\Desktop\Msc IT Practical>
kh/AppData/Local/Programs/Python/Python310/python.exe" "c:/Us
ve/Desktop/Msc IT Practical/ML_Prac_3a.py"
Intercept: 0.2142857142857153
Coefficient: 2.0071428571428567
MSE: 0.01951530612244882
R-squared: 0.9978316326530613
PS C:\Users\Saqlain Shaikh\OneDrive\Desktop\Msc IT Practical>
```

Practical 3B**AIM:**

- **Multiple Linear Regression** Extend linear regression to multiple features. Handle feature selection and potential multicollinearity.

CODE:

```
import pandas as pd
from sklearn.model_selection import train_test_split
from sklearn.linear_model import LinearRegression
from sklearn.metrics import mean_squared_error, r2_score
from statsmodels.stats.outliers_influence import
variance_inflation_factor

data = {
    'Feature1': [1, 2, 3, 4, 5, 6, 7, 8, 9, 10],
    'Feature2': [2, 4, 6, 8, 10, 12, 14, 16, 18, 20],
    'Feature3': [1, 3, 2, 4, 3, 5, 6, 7, 8, 9],
    'Target': [3, 7, 5, 9, 11, 15, 17, 21, 23, 27]
}
df = pd.DataFrame(data)

X = df[['Feature1', 'Feature2', 'Feature3']]
Y = df['Target']

vif_data = pd.DataFrame()
vif_data['Feature'] = X.columns
vif_data['VIF'] = [variance_inflation_factor(X.values, i) for i in
range(X.shape[1])]

print("\nVIF for Features:")
print(vif_data)

X_train, X_test, Y_train, Y_test = train_test_split(X, Y,
test_size=0.2,
random_state=42)

model = LinearRegression()
model.fit(X_train, Y_train)

print("\nIntercept:", model.intercept_)
print("Coefficients:", model.coef_)

Y_pred = model.predict(X_test)

print("MSE:", mean_squared_error(Y_test, Y_pred))
print("R-squared:", r2_score(Y_test, Y_pred))
```

```

double vif = 1. / (1. - r_squared_i)

VIF for Features:
  Feature      VIF
0 Feature1      inf
1 Feature2      inf
2 Feature3 69.271726

Intercept: -1.4322344322344343
Coefficients: [0.33846154 0.67692308 1.21611722]
MSE: 1.1180882609454033
R-squared: 0.982529870922728
PS C:\Users\Saqlain Shaikh\OneDrive\Desktop\Msc IT Practical>

```

OUTPUT:

```

Ridge: [ 2.5807138 -1.77414379 0.90149729] 0.04458190346459912 0.9584098704060822
Lasso: [ 1.60397718 -0.98933401 0.          ] 0.31614295332261333 0.705072565837569
ElasticNet: [ 1.38765543 -0.98031813 0.22377714] 0.34657204388252716 0.67668549122966
67
PS C:\Users\Saqlain Shaikh\OneDrive\Desktop\Msc IT Practical>

```

PRACTICAL3C**AIM:**

- Potential multicollinearity. Regularized Linear Models (Ridge, Lasso, Elastic Net)
- Implement regression variants like LASSO and Ridge on any generated dataset. 4 Discriminative Models OC2,0

CODE:

```
import numpy as np
from sklearn.model_selection import train_test_split
from sklearn.linear_model import Ridge, Lasso, ElasticNet
from sklearn.metrics import mean_squared_error, r2_score
np.random.seed(42)
X = np.random.rand(100, 3)
Y = 3 * X[:, 0] - 2 * X[:, 1] + X[:, 2] + np.random.normal(0, 0.1, 100)
X_train, X_test, Y_train, Y_test = train_test_split(X, Y,
test_size=0.2,
random_state=42)
ridge_model = Ridge(alpha=1.0)
ridge_model.fit(X_train, Y_train)
ridge_pred = ridge_model.predict(X_test)
print("Ridge:", ridge_model.coef_, mean_squared_error(Y_test,
ridge_pred),
r2_score(Y_test, ridge_pred))
lasso_model = Lasso(alpha=0.1)
lasso_model.fit(X_train, Y_train)
lasso_pred = lasso_model.predict(X_test)
print("Lasso:", lasso_model.coef_, mean_squared_error(Y_test,
lasso_pred),
r2_score(Y_test, lasso_pred))
elasticnet_model = ElasticNet(alpha=0.1, l1_ratio=0.5)
elasticnet_model.fit(X_train, Y_train)
elasticnet_pred = elasticnet_model.predict(X_test)
print("ElasticNet:", elasticnet_model.coef_,
mean_squared_error(Y_test,
elasticnet_pred), r2_score(Y_test, elasticnet_pred))
```

OUTPUT:

Aim: Discriminative Models

[illegible]

PRACTICAL 4a:

- Perform binary classification using logistic regression. Calculate accuracy, precision, recall, and understand the ROC curve.

```
import numpy as np
import pandas as pd
from sklearn.model_selection import train_test_split
from sklearn.linear_model import LogisticRegression
from sklearn.metrics import (
    accuracy_score, precision_score, recall_score,
    roc_curve, auc
)
import matplotlib.pyplot as plt

np.random.seed(42)
X = np.random.rand(100, 2)
Y = (X[:, 0] + X[:, 1] > 1).astype(int)

X_train, X_test, Y_train, Y_test = train_test_split(X, Y, test_size=0.3,
    random_state=42)

model = LogisticRegression()
model.fit(X_train, Y_train)

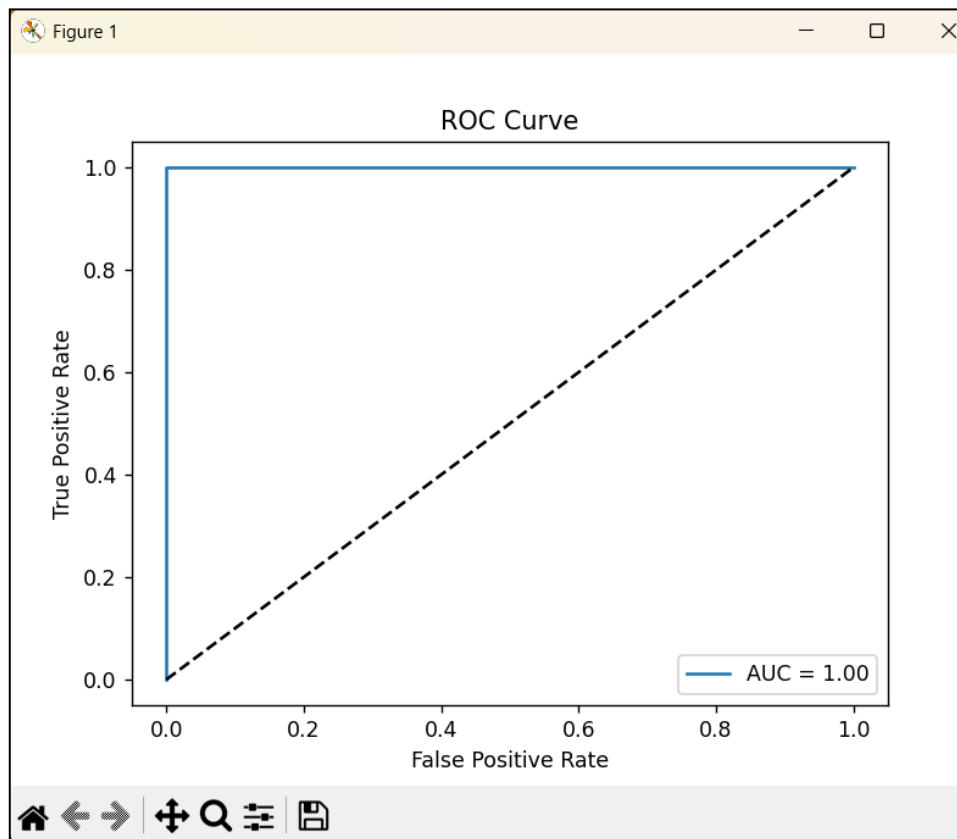
Y_pred = model.predict(X_test)
Y_pred_prob = model.predict_proba(X_test)[:, 1]

print("Accuracy:", accuracy_score(Y_test, Y_pred))
print("Precision:", precision_score(Y_test, Y_pred))
print("Recall:", recall_score(Y_test, Y_pred))

fpr, tpr, _ = roc_curve(Y_test, Y_pred_prob)
plt.plot(fpr, tpr, label=f"AUC = {auc(fpr, tpr):.2f}")
plt.plot([0, 1], [0, 1], 'k--')
plt.xlabel("False Positive Rate")
plt.ylabel("True Positive Rate")
plt.title("ROC Curve")
plt.legend(loc="lower right")
plt.show()
```

Output:

```
Accuracy: 0.8666666666666667
Precision: 1.0
Recall: 0.7333333333333333
```



PRACTICAL 4b:

- Implement and demonstrate k-nearest Neighbor algorithm. Read the training data from a CSV file and build the model to classify a test sample. Print both correct and wrong predictions.

Code:

```
import pandas as pd
import numpy as np
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import StandardScaler
from sklearn.neighbors import KNeighborsClassifier
from sklearn.metrics import classification_report, confusion_matrix
import urllib.request
import os
url = "https://archive.ics.uci.edu/ml/machine-learning-
databases/iris/iris.data"
dataset_file = "iris.data"
if not os.path.exists(dataset_file):
    urllib.request.urlretrieve(url, dataset_file)
columns = ["sepal_length", "sepal_width", "petal_length", "petal_width",
"class"]
data = pd.read_csv(dataset_file, header=None, names=columns)
X = data.iloc[:, :-1].values
y = data.iloc[:, -1].values
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2,
random_state=42)
scaler = StandardScaler()
X_train = scaler.fit_transform(X_train)
X_test = scaler.transform(X_test)
k = 3
knn = KNeighborsClassifier(n_neighbors=k)
knn.fit(X_train, y_train)
y_pred = knn.predict(X_test)
conf_matrix = confusion_matrix(y_test, y_pred)
report = classification_report(y_test, y_pred)
print("Confusion Matrix:")
print(conf_matrix)
print("\nClassification Report:")
print(report)
correct_predictions = np.sum(y_test == y_pred)
incorrect_predictions = np.sum(y_test != y_pred)
print(f"\nCorrect Predictions: {correct_predictions}")
print(f"Incorrect Predictions: {incorrect_predictions}")
results = pd.DataFrame({"Actual": y_test, "Predicted": y_pred})
print("\nSample Results:")
print(results.head())
```

Output:

```
Confusion Matrix:
```

```
[[10  0  0]
 [ 0  9  0]
 [ 0  0 11]]
```

```
Classification Report:
```

	precision	recall	f1-score	support
Iris-setosa	1.00	1.00	1.00	10
Iris-versicolor	1.00	1.00	1.00	9
Iris-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

```
Correct Predictions: 30
```

```
Incorrect Predictions: 0
```

```
Sample Results:
```

	Actual	Predicted
0	Iris-versicolor	Iris-versicolor
1	Iris-setosa	Iris-setosa
2	Iris-virginica	Iris-virginica
3	Iris-versicolor	Iris-versicolor
4	Iris-versicolor	Iris-versicolor

Practical4c:

- Build a decision tree classifier or regressor. Control hyper parameters like tree depth to avoid overfitting. Visualize the tree.

Code:

```
import numpy as np
import pandas as pd
from sklearn.datasets import load_iris
from sklearn.model_selection import train_test_split
from sklearn.tree import DecisionTreeClassifier, plot_tree
from sklearn.metrics import accuracy_score
import matplotlib.pyplot as plt

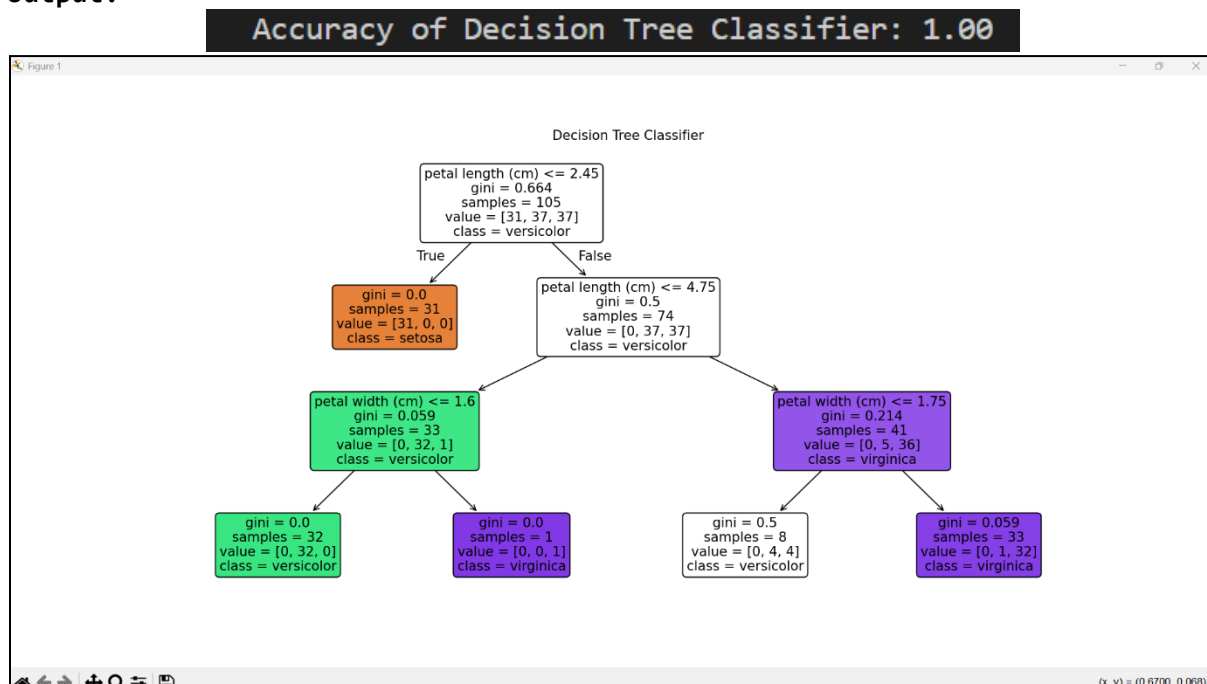
data = load_iris()
X = data.data
y = data.target

X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3,
random_state=42)

clf = DecisionTreeClassifier(max_depth=3, random_state=42)
clf.fit(X_train, y_train)

y_pred = clf.predict(X_test)
accuracy = accuracy_score(y_test, y_pred)
print(f'Accuracy of Decision Tree Classifier: {accuracy:.2f}')
```

```
plt.figure(figsize=(12,8))
plot_tree(clf, filled=True, feature_names=data.feature_names,
class_names=data.target_names, rounded=True)
plt.title("Decision Tree Classifier")
plt.show()
```

Output:

Practical 4d:

- Implement a Support Vector Machine for any relevant dataset.

Code:

```
import numpy as np
import pandas as pd
from sklearn.datasets import load_iris
from sklearn.model_selection import train_test_split
from sklearn.svm import SVC
from sklearn.metrics import accuracy_score
import matplotlib.pyplot as plt
from sklearn.decomposition import PCA

data = load_iris()
X = data.data
y = data.target

X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3,
random_state=42)

svm_clf = SVC(kernel='linear', random_state=42)
svm_clf.fit(X_train, y_train)

y_pred = svm_clf.predict(X_test)
accuracy = accuracy_score(y_test, y_pred)
print(f'Accuracy of SVM classifier: {accuracy:.2f}')
```



```
pca = PCA(n_components=2)
X_pca = pca.fit_transform(X)

svm_clf_2d = SVC(kernel='linear', random_state=42)
svm_clf_2d.fit(X_pca, y)

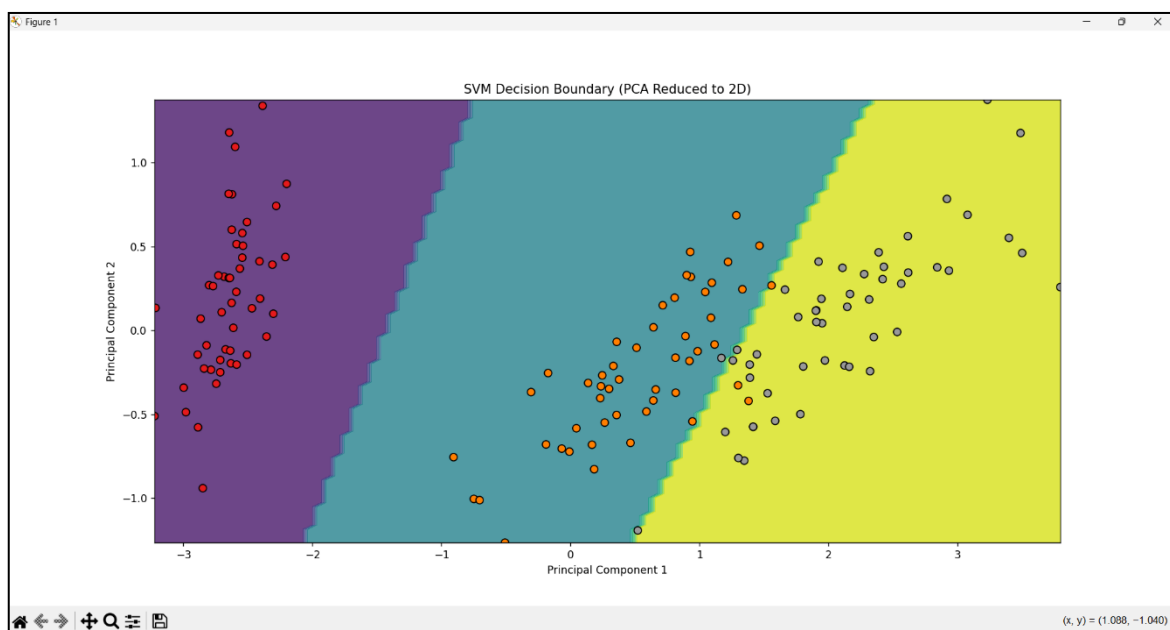
xx, yy = np.meshgrid(np.linspace(X_pca[:, 0].min(), X_pca[:, 0].max(),
100),
                    np.linspace(X_pca[:, 1].min(), X_pca[:, 1].max(),
100))
Z = svm_clf_2d.predict(np.c_[xx.ravel(), yy.ravel()])
Z = Z.reshape(xx.shape)

plt.contourf(xx, yy, Z, alpha=0.8)
plt.scatter(X_pca[:, 0], X_pca[:, 1], c=y, edgecolors='k', marker='o',
s=50,
cmap=plt.cm.Set1)
plt.title("SVM Decision Boundary (PCA Reduced to 2D)")
plt.xlabel("Principal Component 1")
plt.ylabel("Principal Component 2")
plt.show()
```

Output:

PROBLEMS OUTPUT DEBUG CONSOLE TERMINAL PORTS

```
PS C:\Users\Shivang Singh> python -u "d:\6482_Shivang\MLPRACTICAL\4d prac.py"  
Accuracy of SVM classifier: 1.00
```



Practical4e:

- Train a random forest ensemble. Experiment with the number of trees and feature sampling. Compare performance to a single decision tree.

Code:

```
import numpy as np
import pandas as pd
from sklearn.datasets import load_iris
from sklearn.model_selection import train_test_split
from sklearn.tree import DecisionTreeClassifier
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import accuracy_score

data = load_iris()
X = data.data
y = data.target

X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3,
random_state=42)

dt_clf = DecisionTreeClassifier(random_state=42)
dt_clf.fit(X_train, y_train)
dt_y_pred = dt_clf.predict(X_test)
dt_accuracy = accuracy_score(y_test, dt_y_pred)

rf_clf = RandomForestClassifier(n_estimators=100, max_features='sqrt',
random_state=42)
rf_clf.fit(X_train, y_train)
rf_y_pred = rf_clf.predict(X_test)
rf_accuracy = accuracy_score(y_test, rf_y_pred)

rf_clf_50 = RandomForestClassifier(n_estimators=50, max_features='sqrt',
random_state=42)
rf_clf_50.fit(X_train, y_train)
rf_50_y_pred = rf_clf_50.predict(X_test)
rf_50_accuracy = accuracy_score(y_test, rf_50_y_pred)

print(f"Decision Tree Accuracy: {dt_accuracy:.2f}")
print(f"Random Forest (100 trees) Accuracy: {rf_accuracy:.2f}")
print(f"Random Forest (50 trees) Accuracy: {rf_50_accuracy:.2f}")
```

Output:

```
Decision Tree Accuracy: 1.00
Random Forest (100 trees) Accuracy: 1.00
Random Forest (50 trees) Accuracy: 1.00
PS C:\Users\Shivang Singh> █
```

Practical4f

- Implement a gradient boosting machine (e.g., XGBoost). Tune hyperparameters and explore feature importance.

Code:

```
import numpy as np
import pandas as pd
from sklearn.datasets import load_iris
from sklearn.model_selection import train_test_split
import xgboost as xgb
from sklearn.metrics import accuracy_score
import matplotlib.pyplot as plt

data = load_iris()
X = data.data
y = data.target

X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3,
random_state=42)

params = {
    'objective': 'multi:softmax',
    'num_class': 3,
    'max_depth': 3,
    'learning_rate': 0.1,
    'n_estimators': 100,
    'subsample': 0.8,
    'colsample_bytree': 0.8,
    'eval_metric': 'merror'
}

xgb_model = xgb.XGBClassifier(**params)
xgb_model.fit(X_train, y_train)

y_pred = xgb_model.predict(X_test)

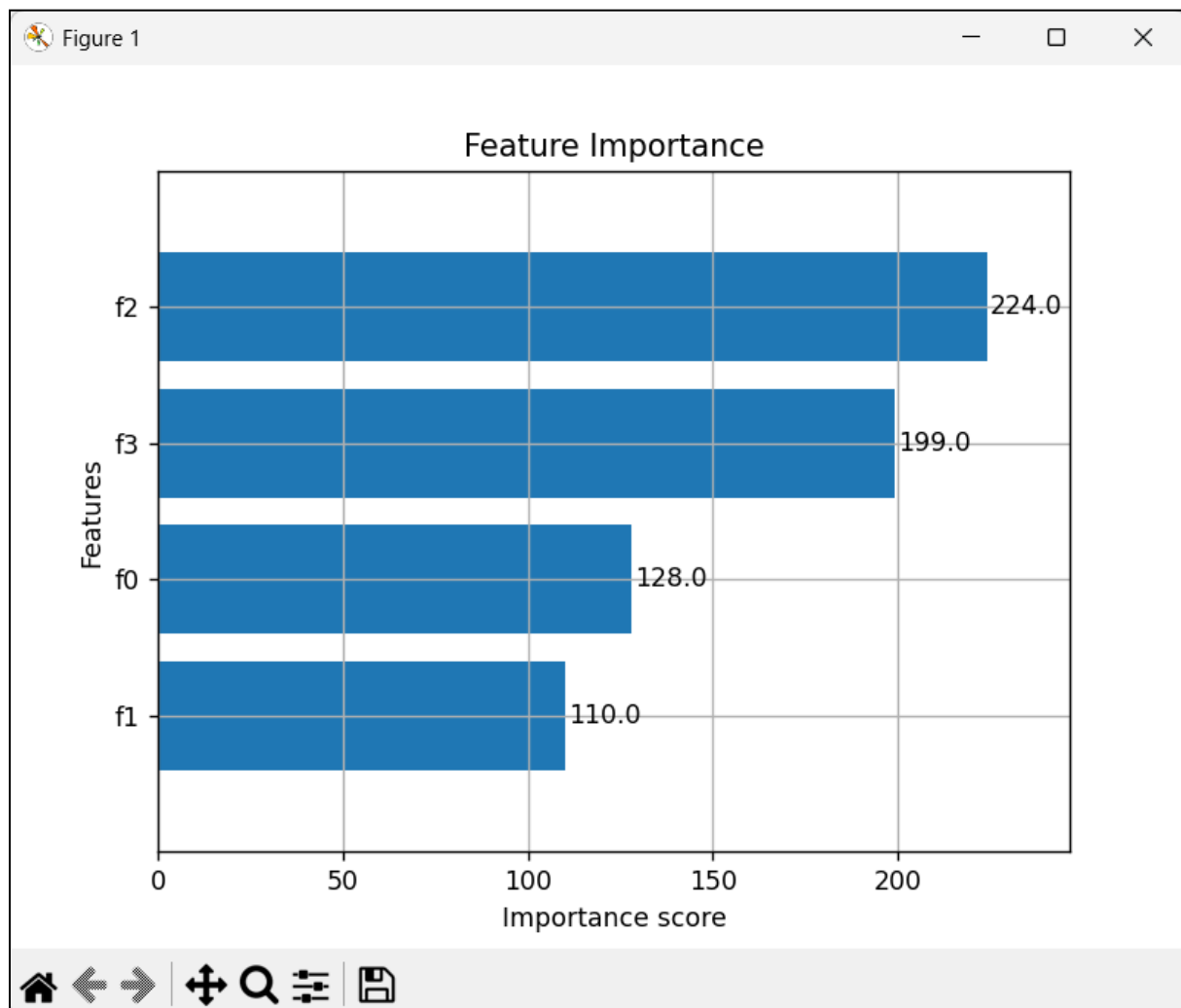
accuracy = accuracy_score(y_test, y_pred)
print(f'Accuracy of XGBoost model: {accuracy:.2f}')
```



```
xgb.plot_importance(xgb_model, importance_type='weight',
max_num_features=4,
height=0.8)
plt.title('Feature Importance')
plt.show()
```

Output:

Accuracy of XGBoost model: 1.00



Practical No: - 5

Aim: Generative Models

Writeups:

[illegible]

Practical5a:

- Implement and demonstrate the working of a Naive Bayesian classifier using a sample data set. Build the model to classify a test sample.

Code:

```
import numpy as np
import pandas as pd
from sklearn.datasets import load_iris
from sklearn.model_selection import train_test_split
from sklearn.naive_bayes import GaussianNB
from sklearn.metrics import accuracy_score
data = load_iris()
X = data.data
y = data.target
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3,
random_state=42)
nb_classifier = GaussianNB()
nb_classifier.fit(X_train, y_train)
y_pred = nb_classifier.predict(X_test)
accuracy = accuracy_score(y_test, y_pred)
print(f'Accuracy of Naive Bayes classifier: {accuracy:.2f}')
test_sample = X_test[0].reshape(1, -1)
predicted_class = nb_classifier.predict(test_sample)
print(f'Predicted class for the test sample: {predicted_class[0]}')
```

Output:

```
===== RESTART: C:/Users/hp/Downloads/ML/ML
Accuracy of Naive Bayes classifier: 0.98
Predicted class for the test sample: 1
> |
```

Practical 5b:

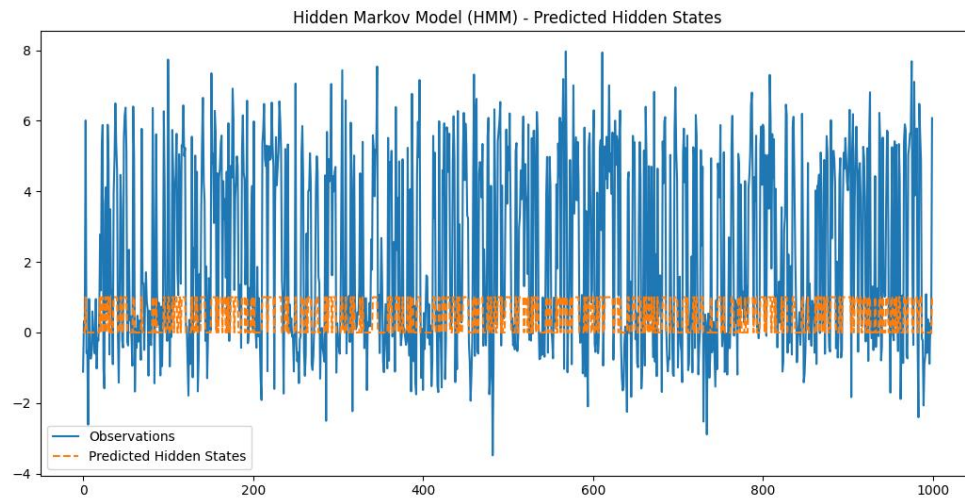
- Implement Hidden Markov Models using hmmlearn

Code:

```
import numpy as np
from hmmlearn.hmm import GaussianHMM
import matplotlib.pyplot as plt
np.random.seed(42)
hidden_states = 2
n_samples = 1000
trans_probs = np.array([[0.7, 0.3], [0.4, 0.6]])
means = np.array([[0.0], [5.0]])
# Covariance matrix needs to be 3D with shape (n_components, n_dim, n_dim)
covars = np.array([[[1.0]], [[1.0]]]) # Correct shape for 'full'
covariance
type
model = GaussianHMM(n_components=hidden_states, covariance_type="full",
n_iter=1000)
model.startprob_ = np.array([0.6, 0.4])
model.transmat_ = trans_probs
model.means_ = means
model.covars_ = covars
X, Z = model.sample(n_samples)

model.fit(X)
```

```
predicted_states = model.predict(X)
plt.figure(figsize=(12, 6))
plt.plot(X, label='Observations')
plt.plot(predicted_states, label='Predicted Hidden States', linestyle='--')
plt.legend()
plt.title('Hidden Markov Model (HMM) - Predicted Hidden States')
plt.show()
```



Practical No: - 6

Aim: Probabilistic Models

Writeups:

[illegible]

Practical6a

- Implement Bayesian Linear Regression to explore prior and posterior distribution.

Code:

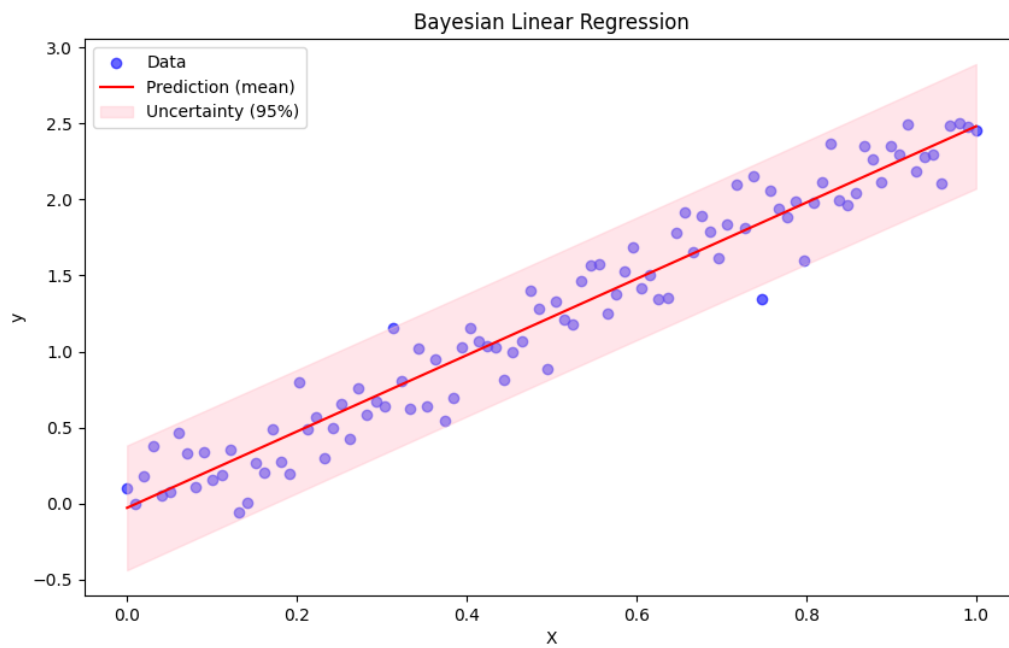
```
import numpy as np
import matplotlib.pyplot as plt
from sklearn.model_selection import train_test_split
np.random.seed(42)
X = np.linspace(0, 1, 100).reshape(-1, 1)
y = 2.5 * X.squeeze() + np.random.normal(0, 0.2, X.shape[0])
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2,
random_state=42)
class BayesianLinearRegression:
    def __init__(self, alpha=1.0, beta=1.0):
        self.alpha = alpha
        self.beta = beta
        self.w_mean = None
        self.w_cov = None
    def fit(self, X, y):
        X = np.hstack((np.ones((X.shape[0], 1)), X))
        S0_inv = self.alpha * np.eye(X.shape[1])
        SN_inv = S0_inv + self.beta * X.T @ X
        SN = np.linalg.inv(SN_inv)
        mN = self.beta * SN @ X.T @ y
        self.w_mean = mN
        self.w_cov = SN

    def predict(self, X, return_std=False):
        X = np.hstack((np.ones((X.shape[0], 1)), X))
        mean = X @ self.w_mean
        if return_std:
            variance = 1 / self.beta + np.sum(X @ self.w_cov * X, axis=1)
            return mean, np.sqrt(variance)
        return mean

blr = BayesianLinearRegression(alpha=1.0, beta=25.0)
blr.fit(X_train, y_train)

X_pred = np.linspace(0, 1, 100).reshape(-1, 1)
y_pred, y_std = blr.predict(X_pred, return_std=True)
plt.figure(figsize=(10, 6))
plt.scatter(X, y, label="Data", color="blue", alpha=0.6)
plt.plot(X_pred, y_pred, label="Prediction (mean)", color="red")
plt.fill_between(
    X_pred.squeeze(),
    y_pred - 2 * y_std,
    y_pred + 2 * y_std,
    color="pink",
    alpha=0.4,
    label="Uncertainty (95%)", )
plt.title("Bayesian Linear Regression")
plt.xlabel("X")
plt.ylabel("y")
plt.legend()
plt.show()
```

Output:

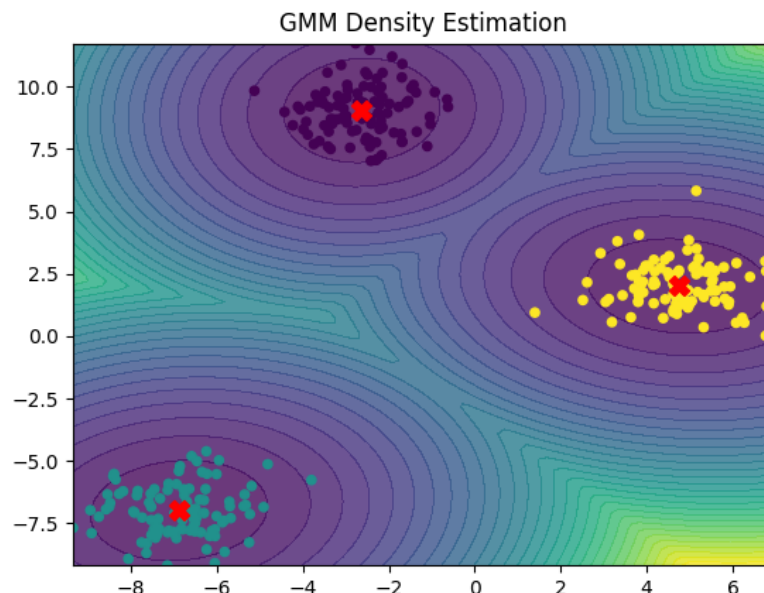


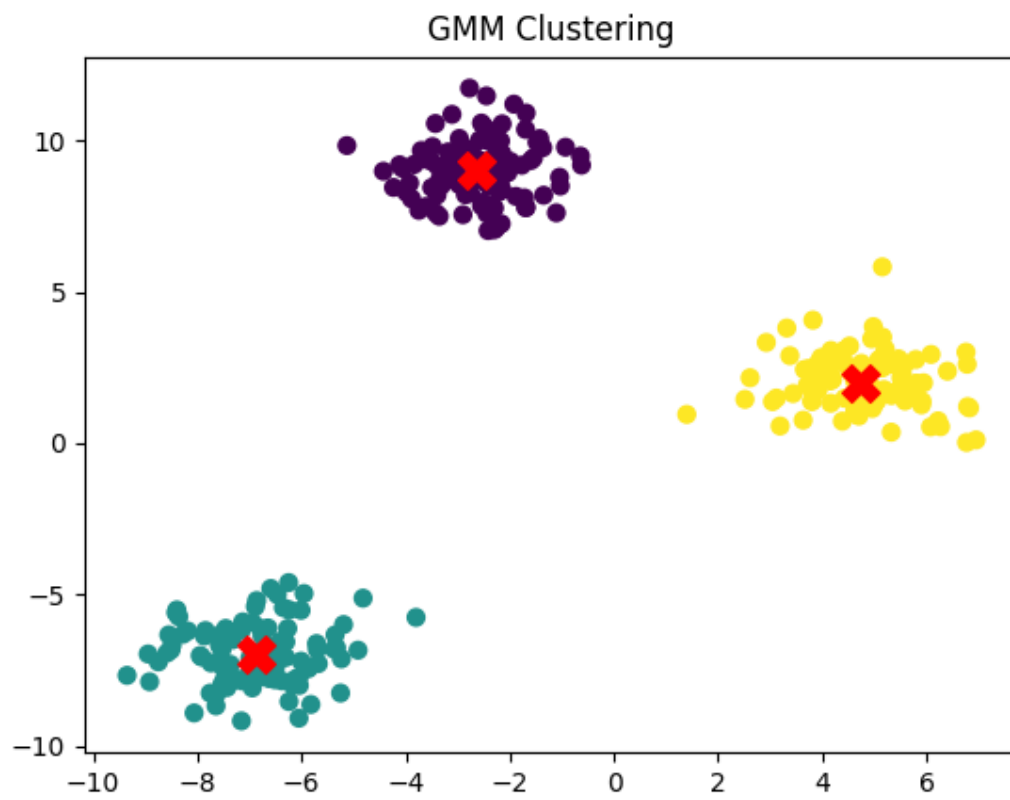
Practical6b

- Implement Gaussian Mixture Models for density estimation and unsupervised clustering

Code:

```
import numpy as np
from sklearn.mixture import GaussianMixture
from sklearn.datasets import make_blobs
import matplotlib.pyplot as plt
X, _ = make_blobs(n_samples=300, centers=3, cluster_std=1.0,
random_state=42)
gmm = GaussianMixture(n_components=3, covariance_type='full',
random_state=42).fit(X)
labels = gmm.predict(X)
plt.scatter(X[:, 0], X[:, 1], c=labels, cmap='viridis', s=40)
plt.scatter(gmm.means_[:, 0], gmm.means_[:, 1], c='red', s=200, marker='X')
plt.title("GMM Clustering")
plt.show()
x, y = np.linspace(X[:, 0].min(), X[:, 0].max(), 100), np.linspace(X[:,
1].min(), X[:, 1].max(), 100)
X_grid, Y_grid = np.meshgrid(x, y)
grid_points = np.c_[X_grid.ravel(), Y_grid.ravel()]
Z = -gmm.score_samples(grid_points).reshape(X_grid.shape)
plt.contourf(X_grid, Y_grid, Z, levels=30, cmap='viridis', alpha=0.8)
plt.scatter(X[:, 0], X[:, 1], c=labels, cmap='viridis', s=20)
plt.scatter(gmm.means_[:, 0], gmm.means_[:, 1], c='red', s=100, marker='X')
plt.title("GMM Density Estimation")
plt.show()
```

Output:



Practical No: - 7

Aim: Model Evaluation and Hyperparameter Tuning.

Writeups:

[illegible]

Practical 7a:

- Implement cross-validation techniques (k-fold, stratified, etc.) for robust model evaluation.

Code:

```
import numpy as np
from sklearn.datasets import make_classification
from sklearn.model_selection import KFold, StratifiedKFold, cross_val_score
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import accuracy_score
X, y = make_classification(n_samples=1000, n_features=10, n_informative=5,
n_redundant=2, random_state=42)
model = RandomForestClassifier(random_state=42)
kf = KFold(n_splits=5, shuffle=True, random_state=42)
kfold_scores = cross_val_score(model, X, y, cv=kf, scoring='accuracy')
print("K-Fold Cross-Validation Scores:", kfold_scores)
print("K-Fold Average Accuracy:", np.mean(kfold_scores))
skf = StratifiedKFold(n_splits=5, shuffle=True, random_state=42)
stratified_scores = cross_val_score(model, X, y, cv=skf,
scoring='accuracy')
print("Stratified K-Fold Cross-Validation Scores:", stratified_scores)
print("Stratified K-Fold Average Accuracy:", np.mean(stratified_scores))
from sklearn.model_selection import train_test_split
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3,
random_state=42)
model.fit(X_train, y_train)
y_pred = model.predict(X_test)
holdout_accuracy = accuracy_score(y_test, y_pred)
print("Holdout Validation Accuracy:", holdout_accuracy)
```

Output:

```
= RESTART: C:/Users/hp/OneDrive/Documents/MSCIT/PART 2/SEMESTER 3/ML/ML p7a.py
K-Fold Cross-Validation Scores: [0.955 0.915 0.9  0.925 0.935]
K-Fold Average Accuracy: 0.9260000000000002
Stratified K-Fold Cross-Validation Scores: [0.925 0.955 0.94  0.925 0.94 ]
Stratified K-Fold Average Accuracy: 0.937
Holdout Validation Accuracy: 0.94
```

Practical 7b:

- **Systematically explore combinations of hyperparameters to optimize model performance. (use grid and randomized search)**

Code:

```
import numpy as np
from sklearn.datasets import make_classification
from sklearn.ensemble import RandomForestClassifier
from sklearn.model_selection import GridSearchCV, RandomizedSearchCV
from sklearn.model_selection import train_test_split
from sklearn.metrics import accuracy_score
X, y = make_classification(n_samples=1000, n_features=10, n_informative=5,
n_redundant=2, random_state=42)
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3,
random_state=42)
model = RandomForestClassifier(random_state=42)
param_grid = {
    'n_estimators': [50, 100, 200],
    'max_depth': [None, 10, 20, 30],
    'min_samples_split': [2, 5, 10],
    'min_samples_leaf': [1, 2, 4]
}
grid_search = GridSearchCV(estimator=model, param_grid=param_grid, cv=5,
n_jobs=-1, verbose=2, scoring='accuracy')
grid_search.fit(X_train, y_train)
print("Best parameters from Grid Search:", grid_search.best_params_)
best_model_grid = grid_search.best_estimator_
y_pred_grid = best_model_grid.predict(X_test)
grid_accuracy = accuracy_score(y_test, y_pred_grid)

print("Grid Search Model Accuracy:", grid_accuracy)
param_dist = {
    'n_estimators': [50, 100, 200, 300],
    'max_depth': [None, 10, 20, 30, 40],
    'min_samples_split': [2, 5, 10, 15],
    'min_samples_leaf': [1, 2, 4, 6]
}
random_search = RandomizedSearchCV(estimator=model,
param_distributions=param_dist, n_iter=10, cv=5, n_jobs=-1, verbose=2,
scoring='accuracy', random_state=42)
random_search.fit(X_train, y_train)
print("Best parameters from Randomized Search:", random_search.best_params_)
best_model_random = random_search.best_estimator_
y_pred_random = best_model_random.predict(X_test)
random_accuracy = accuracy_score(y_test, y_pred_random)
print("Randomized Search Model Accuracy:", random_accuracy)
```

Output:

```
= RESTART: C:/Users/hp/OneDrive/Documents/MSCIT/PART 2/SEMESTER 3/ML/ML p7b.py =
Fitting 5 folds for each of 108 candidates, totalling 540 fits
Best parameters from Grid Search: {'max_depth': None, 'min_samples_leaf': 1, 'min_samples_split': 2, 'n_estimators': 200}
Grid Search Model Accuracy: 0.94
Fitting 5 folds for each of 10 candidates, totalling 50 fits
Best parameters from Randomized Search: {'n_estimators': 100, 'min_samples_split': 10, 'min_samples_leaf': 1, 'max_depth': None}
Randomized Search Model Accuracy: 0.9333333333333333
>>|
```

Practical No: - 8

Aim: Bayesian Learning

Writeups:

[illegible]

Practical 8a:

- Implement Bayesian Learning using inferences

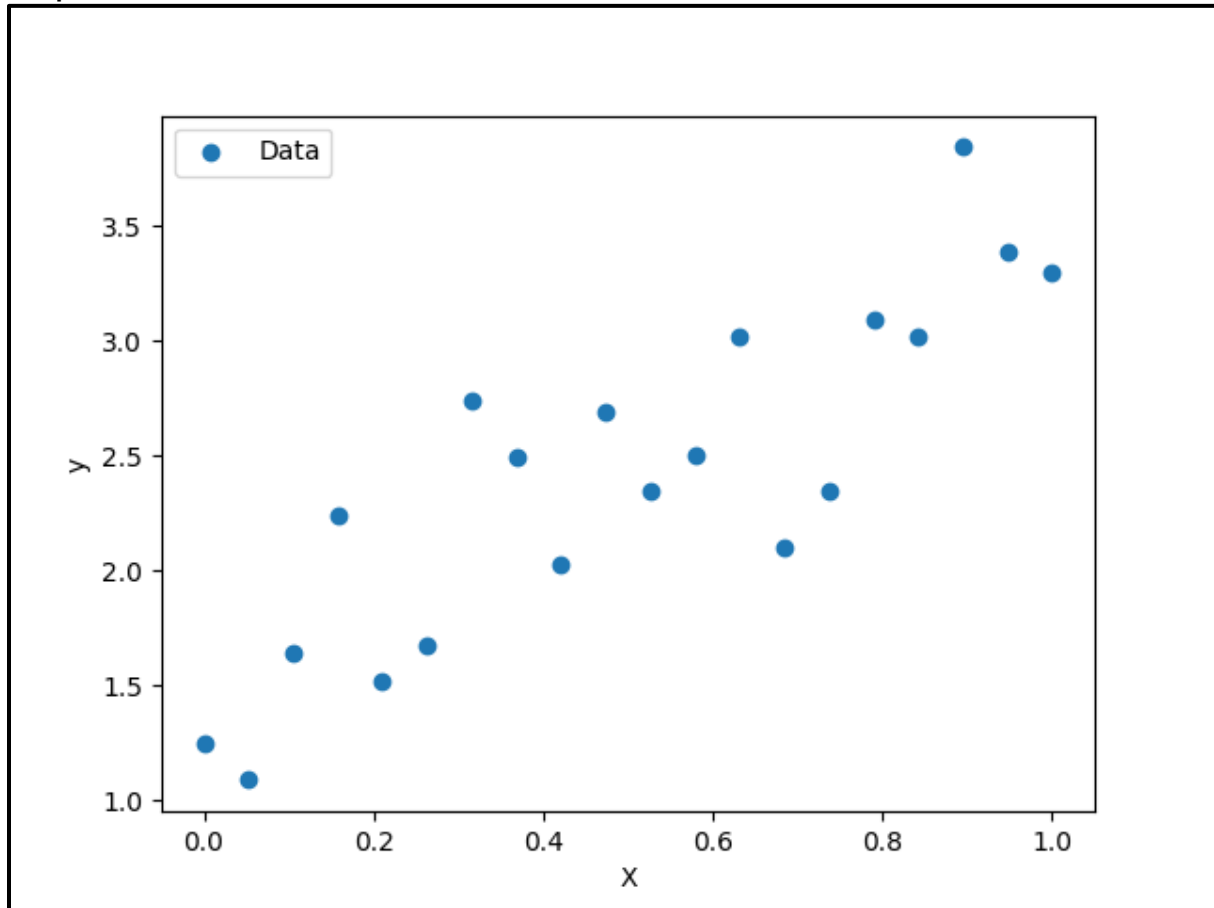
Code:

```
import numpy as np
import matplotlib.pyplot as plt
np.random.seed(42)
X = np.linspace(0, 1, 20)
true_slope = 3
true_intercept = 1
y = true_slope * X + true_intercept + np.random.normal(scale=0.5,
size=X.shape)
plt.scatter(X, y, label="Data")
plt.xlabel("X")
plt.ylabel("y")
plt.legend()
plt.show()
X_ = np.vstack([X, np.ones_like(X)]).T
sigma_prior = 10
mu_prior = np.array([0, 0])
sigma_likelihood = 0.5
sigma_likelihood_inv = np.linalg.inv(sigma_likelihood**2 * np.eye(len(X)))
X_T = X_.T
covariance_post = np.linalg.inv(X_T @ X_ / sigma_likelihood**2 + np.eye(2)
/
sigma_prior**2)
mean_post = covariance_post @ (X_T @ y / sigma_likelihood**2 + mu_prior /
sigma_prior**2)

num_samples = 1000
posterior_samples = np.random.multivariate_normal(mean_post,
covariance_post,
num_samples)
plt.figure(figsize=(12, 6))
plt.subplot(1, 2, 1)
plt.hist(posterior_samples[:, 0], bins=30, color='skyblue',
edgecolor='black')
plt.title("Posterior Distribution of Slope")
plt.subplot(1, 2, 2)
plt.hist(posterior_samples[:, 1], bins=30, color='skyblue',
edgecolor='black')
plt.title("Posterior Distribution of Intercept")
plt.tight_layout()
plt.show()
plt.scatter(X, y, label="Data")
for sample in posterior_samples:
    plt.plot(X, sample[0] * X + sample[1], color='red', alpha=0.05)
plt.xlabel("X")
plt.ylabel("y")
plt.legend()
plt.show()
estimated_slope = mean_post[0]
estimated_intercept = mean_post[1]
```

```
print(f"Estimated Slope: {estimated_slope}")  
print(f"Estimated Intercept: {estimated_intercept}")
```

Output:



Practical 9

Aim: Deep Generative Models

- a) Set up a generator network to produce samples and a discriminator network to distinguish between real and generated data. (Use a simple small dataset)

Goal

We want to build a **GAN** (Generative Adversarial Network) with:

- a **Generator** → makes fake data (from random noise).
- a **Discriminator** → checks if data is real or fake.
- They train together until the generator produces data that looks

real. We'll do this on a **small dataset** (toy 2D data or MNIST digits).

Step-by-Step Instructions:

1. Prepare Python and VS Code

- Make sure Python is installed (type `python --version` in terminal).
- Install **VS Code** and the **Python extension**.

2. Create a project folder

- Open a terminal (cmd, PowerShell, or bash) and type: `mkdir gan-practical`

```
mkdir gan-practical
cd gan-practical
cd gan-practical
```

3. Create a virtual environment (to keep things clean)

```
python -m venv venv
```

```
python -m venv venv
```

Activate it:

- Windows:

```
venv\Scripts\activ
```

Now your terminal should show (venv) in front.

4. Install needed libraries

```
pip install torch torchvision matplotlib numpy tqdm
```

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```

This installs PyTorch and helper tools.

5. Open folder in VS Code

- Open VS Code → **File** → **Open Folder** → select gan-practical.
- In VS Code, press Ctrl+Shift+P → type **Python: Select Interpreter** → choose the one inside venv.

6. Create Python file

- Make a file in VS Code called gan_2d_toy.py.
Paste this small code (toy GAN):

```
import torch, torch.nn as nn
import matplotlib.pyplot
as plt import numpy as np,
os

device = torch.device("cuda" if torch.cuda.is_available() else
"cpu") os.makedirs("samples", exist_ok=True)

# Real data: points around (1,0), (-1,0), (0,1),
(0,-1) def sample_real(batch):
    centers = np.array([[1,0],[-1,0],[0,1],[0,-1]],
dtype=np.float32) idx = np.random.randint(0,4,batch)
    pts = centers[idx] +
    0.1*np.random.randn(batch,2).astype(np.float32) return
    torch.tensor(pts, device=device)

# Generator: noise -> 2D
point class
Generator(nn.Module):
    def __init__(self):
        super().__init__()
        self.net = nn.Sequential(
```



```

# Discriminator: 2D point -> prob class
Discriminator(nn.Module):
    def __init__(self):
        super().__init__()
        self.net = nn.Sequential(
            nn.Linear(2,64), nn.LeakyReLU(0.2), nn.Linear(64,64), nn.LeakyReLU(0.2),
            nn.Linear(64,1), nn.Sigmoid()
        )
    def forward(self,x): return self.net(x)

G, D = Generator().to(device), Discriminator().to(device)
optG = torch.optim.Adam(G.parameters(), lr=2e-4, betas=(0.5,0.999)) optD =
torch.optim.Adam(D.parameters(), lr=2e-4, betas=(0.5,0.999)) bce = nn.BCELoss()

for it in range(1000): # 1. Train D
    real = sample_real(128)
    fake = G(torch.randn(128,2,device=device)).detach()
    lossD = (bce(D(real), torch.ones(128,1,device=device)) +
             bce(D(fake), torch.zeros(128,1,device=device))) * 0.5 optD.zero_grad(); lossD.backward();
    optD.step()

    # 2. Train G
    z = torch.randn(128,2,device=device) fake = G(z)
    lossG = bce(D(fake), torch.ones(128,1,device=device)) optG.zero_grad();
    lossG.backward(); optG.step()

    # Save sample plot every 200 steps if (it+1)%200==0:
    with torch.no_grad():
        gen = G(torch.randn(1000,2,device=device)).cpu().numpy() real =
        sample_real(1000).cpu().numpy()
        plt.scatter(real[:,0],real[:,1],alpha=0.2)
        plt.scatter(gen[:,0],gen[:,1],alpha=0.6)
        plt.savefig(f'samples/iter{it+1}.png'); plt.close() print("Saved sampl at iter",
        it+1)

```

7. Run the code

In VS Code terminal:

☞ This will create a folder samples/ with pictures showing how fake points move closer to real ones.

What is happening?

1. **Generator** starts with random noise → tries to make fake points.
2. **Discriminator** learns to tell apart real vs fake.

3. Both play a “game”:
 - o D gets better at spotting fakes.
 - o G gets better at fooling D.
4. After training, G can produce points similar to the real data.

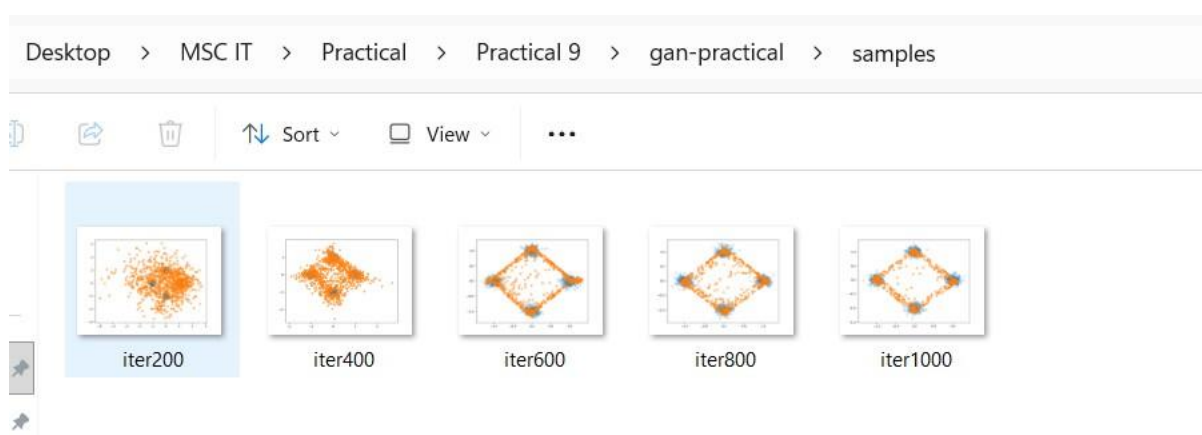
What to submit for practical

- The code (gan_2d_toy.py).
- The sample images (samples/iter200.png, iter400.png, etc.).
- A short explanation: *“Generator makes fake data from noise, Discriminator checks real vs fake, both train together until generator produces realistic data.”*

Output:

```
C:\Users\Sufiya ahmed\OneDrive\Desktop\MSC IT\Practical\Practical 9\gan-practical>python -u "c:\Users\Sufiya ahmed\
Drive\Desktop\MSC IT\Practical\Practical 9\gan-practical\gan_2d_toy.py"
Saved sample at iter 200
Saved sample at iter 400
Saved sample at iter 600
Saved sample at iter 800
Saved sample at iter 1000
```

Name	Date modified	Type	Size
samples	30-09-2025 21:16	File folder	
venv	30-09-2025 21:14	File folder	
gan_2d_toy	30-09-2025 21:16	Python File	3 KB



If you'll get the below error how to fix it:

Error 1:

```
venv\Scripts\activate : File C:\Users\Sufiya
ahmed\OneDrive\Desktop\MSC IT\Practical\Practical 9\gan-
practical\venv\Scripts\Activate.ps1 cannot be loaded
because running scripts is disabled on this system. For more
information, see about_Execution_Policies at
https://go.microsoft.com/fwlink/?LinkID=135170.
At line:1 char:1
+ venv\Scripts\activate
+ ~~~~~
+ CategoryInfo          : SecurityError: (:) [], PSSecurityException
+ FullyQualifiedErrorId : UnauthorizedAccess
```

Quick Fix (without changing policies)

Instead of PowerShell, open **Command Prompt**:

1. In VS Code → Ctrl+Shift+P → type "**Terminal: Select Default Profile**" → pick **Command Prompt**.
2. Open a new terminal (it should say cmd.exe at the top).
3. Run:

```
venv\Scripts\activate.bat
```

☑Now your terminal will show (venv).

