

# MASTER OF SCIENCE IN

## **COMPUTER SCIENCE**

23CSP201: PRINCIPLES OF DATA SCIENCE LAB

**SUBMITTED** 

BY

II SEMESTER MSC
Computer Science Students

#### **SUBMITTED**

TO

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#### 1. Write a python program to

- i. read multiple files from single folder
- ii. read multiple files from multiple folders

```
#program to read multiple files from single folder
import os
path = os.getcwd()
for file in os.listdir(path):
  if file.endswith(".txt"):
    file path = os.path.join(path, file)
    print(file)
    with open(file_path, 'r') as f:
       print(f.read())
Output:
file1.txt
Hello world!!!
file2.txt
Principles of Data Science
file3.txt
Welcome
#program to read multiple files from multiple folders
import os
def read text files from folders(root folder):
  for folder name, subfolders, filenames in os.walk(root folder):
    print("File name :: ",filenames)
    for filename in filenames:
       if filename.endswith('.txt'):
         file_path = os.path.join(folder_name, filename)
         try:
           with open(file_path, 'r') as file:
              print(folder_name)
              print(filename)
```

```
print(file.read())
         except Exception as e:
           print(f"Error reading file {file_path}: {e}")
root_folder = "F:\Sample"
texts = read_text_files_from_folders(root_folder)
Output:
File name :: ['file1.txt', 'file2.txt', 'file3.txt']
F:\Sample\Sample_subfolder1
file1.txt
Hello world!!!
F:\Sample\Sample subfolder1
file2.txt
Principles of Data Science
F:\Sample\Sample_subfolder1
file3.txt
Welcome
File name :: ['file11.txt', 'sample.txt']
F:\Sample\Sample_subfolder3
file11.txt
Matplotlib is an amazing visualization library in Python for 2D plots of arrays.
It was introduced by John Hunter in the year 2002.
Matplotlib consists of several plots like line, bar, scatter, histogram, etc.
F:\Sample\Sample_subfolder3
sample.txt
Python is a dynamic, high-level, free open source, and interpreted programming language. It
```

supports object-oriented programming as well as procedural-oriented programming.

2. Implement the python program to find central tendency (mean, median, and mode) of data, with and without using built-in function on the data.

```
import numpy as np
import statistics
def find_mean(list1):
  total = 0
  for ele in list1:
    total += ele
  mean = total / n
  return mean
def find median(list1):
  list1.sort()
  print("Sorted list elements are :: ",list1)
  if n % 2 == 0:
    median = (list1[n // 2] + list1[n // 2 - 1]) / 2
  else:
    median = list1[n // 2]
  return median
def find_mode(list1):
  unq list = []
  for ele in list1:
    if ele not in unq_list:
       unq_list.append(ele)
  max_count = 0
  mode list = []
  for ele in ung list:
    currentCount = list1.count(ele)
    if currentCount > max_count:
       max count = currentCount
       mode_list = [ele]
    elif currentCount == max count:
       mode_list.append(ele)
  return mode list
```

```
list1 = []
n = int(input("Enter number of elements :: "))
for i in range(n) :
    list1.append(int(input("Enter a number :: ")))

print("List elements are :: ",list1)
print("Mean (built-in) : ", np.mean(list1))
print("Mean (without built-in) : ",find_mean(list1))
print("Median (built-in) : ", np.median(list1))
print("Median (without built-in) : ",find_median(list1))
print("Mode (built-in) : ", statistics.multimode(list1))
print("Mode (without built-in) : ",find_mode(list1))
```

Enter number of elements :: 7

Enter a number :: 1
Enter a number :: 1
Enter a number :: 2
Enter a number :: 5
Enter a number :: 2
Enter a number :: 8
Enter a number :: 4

List elements are :: [1, 1, 2, 5, 2, 8, 4] Mean (built-in) : 3.2857142857142856

Mean (without built-in): 3.2857142857142856

Median (built-in): 2.0

Sorted list elements are :: [1, 1, 2, 2, 4, 5, 8]

Median (without built-in): 2

Mode (built-in): [1, 2]

Mode (without built-in): [1, 2]

3. Implement a program to perform measure of dispersion (range, variance, standard deviation, IQR), with and without using built-in function on the data.

```
import numpy as np
import math
from scipy import stats
def find_range(list1):
  maxElement = minElement = list1[0]
  for ele in list1:
    if ele > maxElement:
      maxElement = ele
    if ele < minElement:
      minElement = ele
  rangeValue = maxElement - minElement
  return rangeValue
def find variance(list1):
  total = 0
  for ele in list1:
    total += ele
  mean = total / len(list1)
  sumValue = 0
  for i in list1:
    sumValue += (i - mean) ** 2
  variance = sumValue / len(list1)
  return variance
def find_sd(list1):
  sd = math.sqrt(find_variance(list1))
  return sd
def percentile midpoint(data, percent):
  sorted_data = sorted(data)
  n = len(sorted data)
  # Calculate the position of the percentile
  k = (n - 1) * percent
```

```
f = int(k) # floor value
  c = k - f # fractional part
  if c == 0:
    # If k is an integer, return the exact value at that position
    return sorted_data[f]
  else:
    # Midpoint interpolation between the two closest ranks
    return (sorted data[f] + sorted data[f + 1]) / 2
def find iqr midpoint(data):
  Q1 = percentile midpoint(data, 0.25) # Calculate the 25th percentile
  Q3 = percentile midpoint(data, 0.75) # Calculate the 75th percentile
  IQR = Q3 - Q1 # Calculate the IQR
  return IQR
list1 = []
n=int(input("Enter the number of elements ::"))
for i in range(n):
  ele = int(input("Enter the elements ::"))
  list1.append(ele)
print("List elements are\n", list1)
print("The range(without built-in):: ", find_range(list1))
maximum = np.max(list1)
minimum = np.min(list1)
range value = maximum - minimum
print("The range(with built-in):: ", range value)
print("The variance(without built-in):: ", find variance(list1))
print("The variance(with built-in):: ", np.var(list1))
print("The standard deviation(without built-in):: ", find sd(list1))
print("The standard deviation(with built-in):: ", np.std(list1))
iqr = find iqr midpoint(list1)
print("Interquartile Range(without built-in function):", iqr)
IQR = stats.iqr(list1, interpolation='midpoint')
print("IQR(with built-in function):", IQR)
```

Enter the number of elements ::7

Enter the elements ::5
Enter the elements ::2
Enter the elements ::7
Enter the elements ::13
Enter the elements ::8
Enter the elements ::6

Enter the elements ::1
List elements are

[5, 2, 7, 13, 8, 6, 1]

The range(without built-in):: 12
The range(with built-in):: 12

The variance(without built-in):: 13.714285714285714
The variance(with built-in):: 13.714285714285714

The standard deviation(without built-in):: 3.7032803990902057
The standard deviation(with built-in):: 3.7032803990902057

Interquartile Range(without built-in function): 4.0

IQR(with built-in function): 4.0

## 4. Write a program to perform text data pre-processing with and without using built-in functions.

#### #With built-in

import pandas as pd import spacy import string import contractions import re import nltk import emoji

from nltk import word\_tokenize from nltk.corpus import stopwords from nltk.stem import WordNetLemmatizer from nltk.stem import PorterStemmer from num2words import num2words

```
nltk.download('punkt')
nltk.download('stopwords')
nltk.download('whitespace')
nltk.download('wordnet')
nltk.download('omw-1.4')
stop_words = set(stopwords.words("english"))
data = pd.read_csv('Training.tsv',sep='\t')
data.head(10)
```

	tweet_id	text	label
0	1382343793341575169	@IrvineWelsh I don't know about you Irvine but	0
1	1377631738692796417	I bet money if i went n took a covid test righ	0
2	1386448010029240326	@JamesMelville My wife received a POSITIVE Cov	0

0	Out of the 180,000+ people who have had the tw	1361342676340211717	3
0	My whole family is sick af and here I am now i	1386757983254765569	4
0	@renfrew1962 @PeakePolly @J_Deliciouso I'm not	1382001700853125122	5
1	Test came back positive, no surprise. I have c	1383272654212272136	6
0	My Pawpaw has been in the hospital a few days	1374479299047084035	7
0	@MattHancock 4 people I know had covid and rec	1354020426620547072	8
1	I'm going to sound like I have lost my marbles	1362671045136809985	9

data['label'].value\_counts()

#### Output:

0 6266

1 1334

Name: label, dtype: int64

```
ps =PorterStemmer()
lemmatiser = WordNetLemmatizer()
english stopwords = stopwords.words('english')
exclude = set(string.punctuation)
def preprocess(text):
  #text=demoji.findall(df['Text'])
  text = contractions.fix(text.lower(), slang=True)
  text = re.sub(r'\d+', lambda x: num2words(int(x.group(0))), text)
  #text= re.sub(r'\d+', '', text)
  text=re.sub(r'$', ", text)
  text= re.sub(r''',", text )
  text=re.sub('<.*?>','',text)
  text=re.sub(r'http\S+', ", text)
  #text=emoji.demojize(text, delimiters=(" ", " "))
  text = ".join(ch for ch in text if ch not in exclude)
  tokens = word tokenize(text)
  #print("Tokens:", tokens)
  text = [t for t in tokens if t not in english stopwords]
```

```
text = " ".join(text)
return text

import emoji
#import demoji
#demoji.download_codes()
def emo(text):
    temp=emoji.demojize(text,delimiters=(" "," "))
    temp=temp.replace("_"," ")
    return temp
data['emo']=data["text"].apply(lambda x:emo(x))
data["clean_text"]=data['emo'].apply(lambda X: preprocess(X))
```

	tweet_id	text	label	emo	clean_text
0	1382343793341575169	@IrvineWelsh I don't know about you Irvine but	0	@IrvineWelsh I don't know about you Irvine but	irvinewelsh know irvine keep told covid exist
1	1377631738692796417	I bet money if i went n took a covid test righ	0	I bet money if i went n took a covid test righ	bet money went n took covid test right going t
2	1386448010029240326	@JamesMelville My wife received a POSITIVE Cov	0	@JamesMelville My wife received a POSITIVE Cov	jamesmelville wife received positive covid tes
3	1361342676340211717	Out of the 180,000+ people who have had the tw	0	Out of the 180,000+ people who have had the tw	one hundred eightyzero people two vaccine shot
4	1386757983254765569	My whole family is sick af and here I am now i	0	My whole family is sick af and here I am now i	whole family sick af hospital heart palpitatio

#### #Without built-in

import pandas as pd
data = pd.read\_csv('Training.tsv',sep='\t')
data

#### **Output:**

	tweet_id	text	label		
0	1382343793341575169	@IrvineWelsh I don't know about you Irvine but	0		
1	1 1377631738692796417 I bet money if i went n took a covid test righ				
2	1386448010029240326	@JamesMelville My wife received a POSITIVE Cov	0		
3	1361342676340211717	Out of the 180,000+ people who have had the tw	0		
4	1386757983254765569	My whole family is sick af and here I am now i	0		
5	1382001700853125122	@renfrew1962 @PeakePolly @J_Deliciouso I'm not	0		
6	1383272654212272136	Test came back positive, no surprise. I have c	1		
7	1374479299047084035	My Pawpaw has been in the hospital a few days	0		
8	1354020426620547072	@MattHancock 4 people I know had covid and rec	0		
9	1362671045136809985	I'm going to sound like I have lost my marbles	1		

#### # Define English stopwords

english\_stopwords = set(['i', 'me', 'my', 'myself', 'we', 'our', 'ours', 'ourselves', 'you', "you're", "you've","you'd", 'yours', 'yours', 'yourself', 'yourselves', 'he', 'him', 'his', 'himself', 'she', "she's", 'her', 'hers', 'herself', 'it', "it's", 'itself', 'they', 'them', 'their', 'such', 'no', 'nor', 'not', 'only', 'own', 'same', 'so', 'than', 'too', 'very', 's', 't', 'can', 'will', 'just', 'don', "don't", 'should', "should've", 'now', 'd', 'll', 'm', 'o', 're', 've', 'wasn', "wasn't", 'weren', "weren't", 'won', "won't", 'wouldn', "wouldn't"])

#### def normalize apostrophes(text):

# Replace different representations of apostrophes with a single consistent representation text = text.replace("'", "'") # Replace curly apostrophe with straight apostrophe return text

```
# Define function to preprocess text
def preprocess_text(text):
  # Lowercasing apostrophe
  text = to_lowercase(text)
  #Normalising
  text = normalize apostrophes(text)
  # Removing Contractions
  text = remove contraction(text)
  # Converting number to words
  text = convert numbers to words(text)
  # Removing URLs
  text = remove urls(text)
  # Removing special characters
  text = remove special characters(text)
 # Tokenization and removing stopwords
  tokens = text.split()
  tokens = [token for token in tokens if token not in english_stopwords]
 # Joining tokens
  text = ' '.join(tokens
  return text
def remove_contraction(text):
 # Define contractions
  contractions = {
    "ain't": "am not / is not / are not / has not / have not",
    "aren't": "are not",
    "can't": "cannot",
    "could've": "could have",
    "couldn't": "could not",
    "didn't": "did not",
    "doesn't": "does not",
    "don't": "do not",
    "hadn't": "had not",
    "hasn't": "has not",
    "haven't": "have not",
     "i'll": "i will",
    "i'll've": "i will have",
```

```
"i'm": "i am",
    "i've": "i have",
    "isn't": "is not",
    "it'll": "it will",
    "it's": "it is / it has",
    "let's": "let us",
     "we've": "we have",
    "weren't": "were not",
    "what'll": "what will",
    "what're": "what are",
    "what's": "what is / what has",
     "you're": "you are",
    "you've": "you have"
  # Expanding contractions
  for contraction, expansion in contractions.items():
    text = text.replace(contraction, expansion)
  return text
# Function to remove convert number to words
def convert_numbers_to_words(text):
# Define a dictionary mapping numeric words to their corresponding words
  num_words = {
    '0': 'zero',
    '1': 'one',
    '2': 'two',
    '3': 'three',
    '4': 'four',
    '5': 'five',
    '6': 'six',
    '7': 'seven',
    '8': 'eight',
    '9': 'nine'
  }
  # Converting numbers to words
  for digit, word in num_words.items():
    text = text.replace(digit, word)
  return text
```

```
def to_lowercase(text):
  lowercase text = "
  for char in text:
    # Check if character is uppercase
    if 'A' <= char <= 'Z':
       # Convert uppercase to lowercase
       lowercase text += chr(ord(char) + 32)
    else:
       lowercase text += char
  return lowercase text
# Function to remove special characters
def remove_special_characters(text):
  # Define special characters
  special_chars = {'!', '"', '#', '$', '%', '&', """, '(', ')', '*', '+', ',', '-', '.', '/', ':', ';', '<', '=',
            '>', '?', '@', '[', '\\', ']', '^', '_', \`', '\{', '\\', '\}', \~'\
  return ".join(char for char in text if char not in special chars)
# Function to remove URLs
def remove urls(text):
  # Split text into words
  words = text.split()
  # Filter out words that do not start with 'http' or 'https'
  filtered words = [word for word in words if not (word.startswith('http://') or
word.startswith('https://'))]
  # Join the filtered words back into a string
  return ''.join(filtered words)
import emoji
#import demoji
#demoji.download codes()
def emo(text):
  temp=emoji.demojize(text,delimiters=(" "," "))
  temp=temp.replace("_"," ")
  return temp
data['emo']=data["text"].apply(lambda x:emo(x))
```

data["clean\_text"]=data['emo'].apply(lambda X: preprocess\_text(X))
data.head()

	tweet_id	text	label	emo	clean_text
0	1382343793341575169	@IrvineWelsh I	0	@IrvineWelsh I	irvinewelsh
		don't know		don't know	know irvine
		about you		about you Irvine	keep told covid
		Irvine but		but	exist
1	1377631738692796417	I bet money if i	0	I bet money if i	bet money
		went n took a		went n took a	went n took
		covid test		covid test righ	covid test right
		righ			imma te
2	1386448010029240326	@JamesMelvill	0	@JamesMelville	jamesmelville
		e My wife		My wife	wife received
		received a		received a	positive covid
		POSITIVE Cov		POSITIVE Cov	tes
3	1361342676340211717	Out of the	0	Out of the	oneeightzeroze
		180,000+		180,000+	rozerozero
		people who		people who	people two
		have had the		have had the	vaccine sh
		tw		tw	
4	1386757983254765569	My whole	0	My whole family	whole family
		family is sick af		is sick af and	sick af hospital
		and here I am		here I am now	heart
		now i		i	palpitatio

## 5. Write a program to perform Numeric data per-processing with and without using built-in functions.

#With built-in import numpy as np import pandas as pd

df = pd.read\_csv("diabetes.csv")
df

### Output:

	Pregna ncies	Glucos e	BloodPressu re	SkinTh icknes s	Insulin	BM I	Diabete sPedigr eeFunct ion	Ag e	Outcom e
0	6.0	148	72.0	35.0	0.0	33. 6	0.627	50	1
1	1.0	85	66.0	29.0	0.0	26. 6	0.351	31	0
2	8.0	183	64.0	0.0	0.0	23. 3	0.672	32	1
3	1.0	89	66.0	23.0	94.0	28. 1	0.167	21	0
4	NaN	137	40.0	35.0	168.0	43. 1	2.288	33	1
						•••			
764	10.0	101	76.0	48.0	180.0	32. 9	0.171	63	0
765	2.0	122	70.0	27.0	0.0	36. 8	0.340	27	0
766	5.0	121	72.0	23.0	112.0	26. 2	0.245	30	0
767	1.0	126	60.0	0.0	0.0	30. 1	0.349	47	1
768	1.0	93	70.0	31.0	0.0	30. 4	0.315	23	0

df.shape

(769, 9)

df.describe()

## Output:

	Pregn ancie s	Glucose	BloodP ressure	SkinT hickn ess	Insuli n	ВМІ	DiabetesPedigr eeFunction	Age	Outc ome
count	768.0 0000 0	769.000 000	768.000 000	768.0 0000 0	768.0 0000 0	769. 0000 00	769.000000	769.0 0000 0	769.0 0000 0
mean	3.846 354	120.847 854	69.1914 06	20.57 4219	79.79 9479	31.9 8556 6	0.471719	33.23 7971	0.348 505
std	3.368 283	31.9780 03	19.1944 30	15.93 7859	115.2 4400 2	7.88 1425	0.331142	11.75 2850	0.476 807
min	0.000	0.00000	0.00000	0.000	0.000	0.00	0.078000	21.00 0000	0.000
25%	1.000 000	99.0000	63.5000 00	0.000	0.000	27.3 0000 0	0.244000	24.00 0000	0.000
50%	3.000	117.000 000	72.0000 00	23.00 0000	30.50 0000	32.0 0000 0	0.371000	29.00 0000	0.000
75%	6.000	140.000 000	80.0000	32.00 0000	127.2 5000 0	36.6 0000 0	0.626000	41.00 0000	1.000
max	17.00 0000	199.000 000	122.000 000	99.00 0000	846.0 0000 0	67.1 0000 0	2.420000	81.00 0000	1.000

```
#seperate features and class label
```

features = df.iloc[:, :-1]
class\_label = df.iloc[:, -1]

```
def find_duplicates(data):
    duplicate_rows = data[data.duplicated()]
    return duplicate_rows
```

```
duplicate_values = find_duplicates(features)
print("Duplicates values:")
print(duplicate_values)
```

**Duplicates values:** 

Pregnancies Glucose BloodPressure SkinThickness Insulin BMI \

6 1.0 85 66.0 29.0 0.0 26.6

#### DiabetesPedigreeFunction Age

6 0.351 31

```
def remove_duplicates(data):
    unique_data = data.drop_duplicates()
    return unique_data
features = remove_duplicates(features)
print("Data after removing duplicates :")
print(features)
```

#### **Output:**

Data after removing duplicates:

Pr	egnancie	es Gluc	ose BloodPi	ressure	SkinThickness	Insulin	BMI \
0	6.0	148	72.0	35.0	0.0 33.6		
1	1.0	85	66.0	29.0	0.0 26.6		
2	8.0	183	64.0	0.0	0.0 23.3		
3	1.0	89	66.0	23.0	94.0 28.1		
4	NaN	137	40.0	35.0	168.0 43.1		
		••					
764	10.0	101	76.0	48.0	180.0 32.9		
765	2.0	122	70.0	27.0	0.0 36.8		
766	5.0	121	72.0	23.0	112.0 26.2		
767	1.0	126	60.0	0.0	0.0 30.1		
768	1.0	93	70.0	31.0	0.0 30.4		

#### DiabetesPedigreeFunction Age

0 0.627 50 1 0.351 31 2 0.672 32

```
3
             0.167 21
4
             2.288 33
             ... ...
764
              0.171 63
765
              0.340 27
766
              0.245 30
767
              0.349 47
768
              0.315 23
def find number of missing values(data):
  missing_values = data.isnull().sum()
  # Filter out columns with missing values
  missing_values = missing_values[missing_values > 0]
  return missing_values
missing values = find number of missing values(features)
# Print columns with missing values and their respective counts
print("Columns with missing values:")
print(missing values)
Output:
Columns with missing values:
Pregnancies 1
BloodPressure 1
SkinThickness 1
```

Insulin

dtype: int64

```
# 1. Handling missing values
def handle missing values(data, strategy='mean'):
  if strategy == 'mean':
    return data.fillna(data.mean())
  elif strategy == 'max':
    return data.fillna(data.max())
  elif strategy == 'min':
    return data.fillna(data.min())
  elif strategy == 'zero':
    return data.fillna(0)
  elif strategy == 'drop':
    return data.dropna()
features = handle missing values(features)
print("Data after handling missing values:")
print(features)
Output:
Data after handling missing values:
  Pregnancies Glucose BloodPressure SkinThickness Insulin BMI \
                 148
                          72.0
0
    6.000000
                                    35.0
                                           0.0 33.6
1
    1.000000
                 85
                         66.0
                                   29.0
                                          0.0 26.6
2
    8.000000
                 183
                          64.0
                                    0.0
                                          0.0 23.3
3
    1.000000
                 89
                         66.0
                                   23.0 94.0 28.1
4
    3.850065
                 137
                          40.0
                                    35.0 168.0 43.1
                                 ... ...
764 10.000000
                  101
                            76.0
                                      48.0 180.0 32.9
765
     2.000000
                  122
                           70.0
                                     27.0
                                           0.0 36.8
     5.000000
                  121
                           72.0
766
                                     23.0 112.0 26.2
767
      1.000000
                  126
                           60.0
                                      0.0
                                            0.0 30.1
      1.000000
                  93
                           70.0
                                     31.0
768
                                            0.0 30.4
```

DiabetesPedigreeFunction Age

0.627 50

0

1	0.351 31	L
2	0.672 32	2
3	0.167 21	L
4	2.288 33	3
764	0.171	63
765	0.340	27
766	0.245	30
767	0.349	47
768	0.315	23

## #Without built-in

import numpy as np import pandas as pd

df = pd.read\_csv("diabetes.csv")
df

	Pregna ncies	Glucos e	BloodPressu re	SkinTh icknes s	Insulin	BM I	Diabete sPedigr eeFunct ion	Ag e	Outcom e
0	6.0	148	72.0	35.0	0.0	33. 6	0.627	50	1
1	1.0	85	66.0	29.0	0.0	26. 6	0.351	31	0
2	8.0	183	64.0	0.0	0.0	23. 3	0.672	32	1
3	1.0	89	66.0	23.0	94.0	28. 1	0.167	21	0
4	NaN	137	40.0	35.0	168.0	43. 1	2.288	33	1
•••			•••						
764	10.0	101	76.0	48.0	180.0	32. 9	0.171	63	0
765	2.0	122	70.0	27.0	0.0	36. 8	0.340	27	0
766	5.0	121	72.0	23.0	112.0	26. 2	0.245	30	0

767	1.0	126	60.0	0.0	0.0	30. 1	0.349	47	1
768	1.0	93	70.0	31.0	0.0	30. 4	0.315	23	0

df.shape

Output:

(770, 9)

df.describe()

	Pregn ancie s	Glucose	BloodP ressure	SkinT hickn ess	Insuli n	вмі	DiabetesPedigr eeFunction	Age	Outc ome
count	768.0 0000 0	769.000 000	768.000 000	768.0 0000 0	768.0 0000 0	769. 0000 00	769.000000	769.0 0000 0	769.0 0000 0
mean	3.846 354	120.847 854	69.1914 06	20.57 4219	79.79 9479	31.9 8556 6	0.471719	33.23 7971	0.348 505
std	3.368 283	31.9780 03	19.1944 30	15.93 7859	115.2 4400 2	7.88 1425	0.331142	11.75 2850	0.476 807
min	0.000	0.00000	0.00000	0.000	0.000	0.00	0.078000	21.00 0000	0.000
25%	1.000	99.0000	63.5000 00	0.000	0.000	27.3 0000 0	0.244000	24.00 0000	0.000
50%	3.000	117.000 000	72.0000 00	23.00 0000	30.50 0000	32.0 0000 0	0.371000	29.00 0000	0.000
75%	6.000	140.000 000	80.0000	32.00 0000	127.2 5000 0	36.6 0000 0	0.626000	41.00 0000	1.000
max	17.00 0000	199.000 000	122.000 000	99.00 0000	846.0 0000 0	67.1 0000 0	2.420000	81.00 0000	1.000 000

```
#seperate features and class label
features = df.iloc[:, :-1]
class_label = df.iloc[:, -1]
import pandas as pd
def find_duplicates(data):
  duplicate rows = []
  seen rows = set() # To track rows that we have already seen
  # Iterate through each row in the DataFrame
  for index, row in data.iterrows():
    # Convert the row to a tuple to make it hashable
    row tuple = tuple(row)
    # Check if this row tuple has already been seen
    if row tuple in seen rows:
      # Append the duplicate row as a Series object
      duplicate_rows.append(row)
    else:
      seen rows.add(row tuple) # Add the row tuple to the set of seen rows
  # Create a DataFrame from the list of duplicate rows
  columns = data.columns
  duplicate df = pd.DataFrame(duplicate rows, columns=columns)
  return duplicate df
duplicate values = find duplicates(features)
print("Duplicates values:")
print(duplicate_values)
Output:
Duplicates values:
 Pregnancies Glucose BloodPressure SkinThickness Insulin BMI \
      6.0 148.0
                      72.0
                                        0.0 33.6
5
                                 35.0
      1.0 85.0
                      66.0
                                        0.0 26.6
                                29.0
 DiabetesPedigreeFunction Age
5
            0.627 50.0
            0.351 31.0
```

```
def remove_duplicates(data):
    seen_rows = set()
    unique_data = []

for index, row in data.iterrows():
    row_tuple = tuple(row)
    if row_tuple not in seen_rows:
        seen_rows.add(row_tuple)
        unique_data.append(row)

# Convert list of rows back to DataFrame
    unique_data_df = pd.DataFrame(unique_data, columns=data.columns)
    return unique_data_df

features = remove_duplicates(features)
print("Data after removing duplicates:")
print(features)
```

### Data after removing duplicates:

Pregnancies Glucose BloodPressure SkinThickness Insulin BMI \

0	6.0 148.0	72.0	35.0 0.0 33.6
1	1.0 85.0	66.0	29.0 0.0 26.6
2	8.0 183.0	64.0	0.0 0.0 23.3
3	1.0 89.0	66.0	23.0 94.0 28.1
4	NaN 137.0	40.0	35.0 168.0 43.1
765	10.0 101.0	76.0	48.0 180.0 32.9
766	2.0 122.0	70.0	27.0 0.0 36.8
767	5.0 121.0	72.0	23.0 112.0 26.2
768	1.0 126.0	60.0	0.0 0.0 30.1
769	1.0 93.0	70.0	31.0 0.0 30.4

#### DiabetesPedigreeFunction Age

0	0.627 50.0
1	0.351 31.0
2	0.672 32.0
3	0.167 21.0

```
4
             2.288 33.0
              ... ...
765
              0.171 63.0
766
              0.340 27.0
767
              0.245 30.0
768
              0.349 47.0
769
               0.315 23.0
def find number of missing values(data):
  # Create an empty dictionary to store column names and their respective counts of missing
values
  missing_values = {}
  # Iterate through each column in the DataFrame
  for column in data.columns:
    # Count the number of missing values in the column
    missing count = sum(1 for value in data[column] if pd.isna(value))
    # If there are missing values in the column, add it to the dictionary
    if missing count > 0:
      missing_values[column] = missing_count
  # Print columns with missing values and their respective counts
  print("Columns with missing values:")
  for column, count in missing_values.items():
    print(f"{column}: {count}")
missing values = find number of missing values(features)
```

**Columns with missing values:** 

Pregnancies: 1 BloodPressure: 1 SkinThickness: 1

Insulin: 1

import pandas as pd

```
import numpy as np
def handle_missing_values(data, strategy='mean'):
  num_cols = data.shape[1] # Number of columns
  filled_data = data.copy() # Create a copy to modify
  if strategy == 'mean':
    # Calculate column means
    col means = [np.mean(data.iloc[:, col]) for col in range(num cols)]
    # Replace NaN values with column means
    for col in range(num cols):
      col mean = col means[col]
      for row in range(len(data)):
        if pd.isna(data.iloc[row, col]):
           filled data.iloc[row, col] = col mean
  elif strategy == 'max':
    # Calculate column max values
    col max = [np.max(data.iloc[:, col]) for col in range(num cols)]
    # Replace NaN values with column max values
    for col in range(num cols):
      col max value = col max[col]
      for row in range(len(data)):
        if pd.isna(data.iloc[row, col]):
           filled data.iloc[row, col] = col max value
  elif strategy == 'min':
    # Calculate column min values
    col min = [np.min(data.iloc[:, col]) for col in range(num cols)]
    # Replace NaN values with column min values
    for col in range(num cols):
      col_min_value = col_min[col]
      for row in range(len(data)):
        if pd.isna(data.iloc[row, col]):
           filled data.iloc[row, col] = col min value
```

```
elif strategy == 'zero':
    # Replace NaN values with 0
    for col in range(num_cols):
      for row in range(len(data)):
         # Check if the value is NaN
         if pd.isna(data.iloc[row, col]):
           # If it is NaN, replace it with 0
           filled_data.iloc[row, col] = 0
  elif strategy == 'drop':
    # Drop rows with NaN values
    filled_data = data.dropna()
  return filled_data
features = handle missing values(features, 'max')
print("Data after handling missing values:")
print(features)
Output:
Data after handling missing values:
```

	Pregnancies	s Glucose	BloodPre	ssure S	SkinThickne	ss Insulin	BMI '	١
0	6.0 1	48.0	72.0	35.0	0.0 33.6			
1	1.0	35.0	66.0	29.0	0.0 26.6			
2	8.0 1	83.0	64.0	0.0	0.0 23.3			
3	1.0	39.0	66.0	23.0	94.0 28.1			
4	<b>17.0</b> 2	137.0	40.0	35.0	168.0 43.	1		
			•••					
76	5 10.0	101.0	76.0	48.0	180.0 32	2.9		
76	5 2.0	122.0	70.0	27.0	0.0 36.8	}		
76	7 5.0	121.0	72.0	23.0	112.0 26	.2		
768	3 1.0	126.0	60.0	0.0	0.0 30.1			
769	9 1.0	93.0	70.0	31.0	0.0 30.4			

DiabetesPedigreeFunction Age

0 0.627 50.0 1 0.351 31.0

2	0.672 32.0
3	0.167 21.0
4	2.288 33.0
765	0.171 63.0
766	0.340 27.0
767	0.245 30.0
768	0.349 47.0
769	0.315 23.0

6.. Write a python program to read and display various kinds of data (image, text, and numeric) saved in different format using various python libraries.

#### **Image**

```
#code to read and display .png file
import cv2
image=cv2.imread('flower.png')
cv2.imshow("image", image)
cv2.waitKey(0)
cv2.destroyAllWindows()
#code to read and display .jpg file
import cv2
image=cv2.imread('dog.jpg')
cv2.imshow("image", image)
cv2.waitKey(0)
cv2.destroyAllWindows()
#code to read and display .gif file
import cv2
def show_gif(file_path):
  cap = cv2.VideoCapture(file_path)
  while True:
    ret, frame = cap.read()
    if not ret:
      break
    cv2.imshow('GIF Viewer', frame)
    if cv2.waitKey(100) \& 0xFF == ord('q'):
      break
  cap.release()
  cv2.destroyAllWindows()
show_gif("moon.gif")
```

#### **Numeric Data**

#code to read and display .csv file import pandas as pd csvdata=pd.read\_csv("headbrain.csv") csvdata

#### Output:

	Gender	Age Range	Head Size(cm^3)	Brain Weight(grams)
0	1.0	1.0	4512	1530.0
1	NaN	1.0	3738	1297.0
2	1.0	1.0	4261	1335.0
3	1.0	1.0	3777	1282.0
4	1.0	1.0	4177	NaN
•••				•••
232	2.0	2.0	3214	1110.0
233	2.0	2.0	3394	1215.0
234	2.0	2.0	3233	1104.0
235	2.0	2.0	3352	1170.0
236	2.0	2.0	3391	1120.0

#code to read and display .tsv file
import pandas as pd
tsvdata=pd.read\_csv("file.tsv", sep = "\t")
tsvdata

	0	50	5	881250949
0	0	172	5	881250949
1	0	133	1	881250949
2	196	242	3	881250949
3	186	302	3	891717742
4	22	377	1	878887116
•••			•••	
99997	880	476	3	880175444
99998	716	204	5	879795543
99999	276	1090	1	874795795
100000	13	225	2	882399156

100001	12	203	3	879959583
--------	----	-----	---	-----------

#code to read and display excel
import pandas as pd
exceldata=pd.read\_excel("exceldata.xlsx", names = ["Number 1" , "Number 2"])
exceldata

#### Output:

	Number 1	Number 2
0	5.5277	9.13020
1	8.5186	13.66200
2	7.0032	11.85400
3	5.8598	6.82330
4	8.3829	11.88600
•••		•••
91	5.8707	7.20290
92	5.3054	1.98690
93	8.2934	0.14454
94	13.3940	9.05510
95	5.4369	0.61705

#### **Text Data**

#code to read and display .txt file
import pandas as pd
txtdata= pd.read\_csv("records.txt", sep=" ")
Txtdata

#### **Output:**

Python is a dynamic, high-level, free open source, and interpreted programming language. It supports object-oriented programming as well as procedural-oriented programming.

```
#code to read and display json file
import json
with open("sample1-json.json", 'r') as f:
    json_ob = json.load(f)
print(json_ob)
```

{'fruit': 'Apple', 'size': 'Large', 'color': 'Red'}

### 7. Write a python program to read and display video and audio data.

```
#Audio
import librosa
from IPython.display import Audio
# Load audio file
audio path = "sample-file-4.wav"
y, sr = librosa.load(audio_path)
# Play audio
Audio(data=y, rate=sr)
#Video
import cv2
# Path to the video file
video_path = "file_example.mp4"
# Open the video file
cap = cv2.VideoCapture(video path)
# Check if the video opened successfully
if not cap.isOpened():
  print("Error: Could not open the video.")
else:
  # Create a flag to track window status
  window_open = True
  # Loop through each frame in the video
  while window_open:
    # Read a frame from the video
    ret, frame = cap.read()
    # If the frame was read successfully
    if ret:
      # Display the frame
      cv2.imshow('Video', frame)
```

```
# Check for the 'q' key to quit
if cv2.waitKey(25) & 0xFF == ord('q'):
    break
else:
    # Break the loop if the video has ended
    break

# Check if the window is still open
if cv2.getWindowProperty('Video', cv2.WND_PROP_VISIBLE) < 1:
    window_open = False

# Release the video capture object and close the window
cap.release()
cv2.destroyAllWindows()</pre>
```

# 8. Write a program to implement a Naive Bayes classifier for sample training dataset. Also plot the confusion matrix to evaluate the classifier's performance.

import matplotlib.pyplot as plt
import pandas as pd
from sklearn.model\_selection import train\_test\_split
from sklearn.preprocessing import StandardScaler
dataset = pd.read\_csv('Social\_Network\_Ads.csv')
dataset

### **Output:**

	User ID	Gender	Age	EstimatedSalar	Purchased
				у	
0	15624510	Male	19	19000	0
1	15810944	Male	35	20000	0
2	15668575	Female	26	43000	0
3	15603246	Female	27	57000	0
4	15804002	Male	19	76000	0
395	15691863	Female	46	41000	1
396	15706071	Male	51	23000	1
397	15654296	Female	50	20000	1
398	15755018	Male	36	33000	0
399	15594041	Female	49	36000	1

X = dataset.iloc[:, [2,3]].values

y = dataset.iloc[:,4].values

X\_train, X\_test, y\_train, y\_test = train\_test\_split(X, y, test\_size=0.25, random\_state=0)

sc = StandardScaler()

X\_train = sc.fit\_transform(X\_train)

X\_test = sc.transform(X\_test)

from sklearn.naive\_bayes import GaussianNB classifier = GaussianNB(priors=[0.4, 0.6], var\_smoothing=1e-9) classifier.fit(X\_train, y\_train)

### **Output:**

GaussianNB(priors=[0.4, 0.6])

```
#changing hyperparameter values
from sklearn.naive_bayes import GaussianNB
# Example hyperparameter values
custom priors = [0.3, 0.7] # Custom priors for classes
custom_var_smoothing = 1e-8 # Custom var_smoothing value
# Initialize Gaussian Naive Bayes classifier with custom hyperparameters
classifier = GaussianNB(priors=custom priors, var smoothing=custom var smoothing)
# Assuming X train and y train are your training data
classifier.fit(X_train, y_train)
Output:
GaussianNB(priors=[0.3, 0.7], var_smoothing=1e-08)
y_pred = classifier.predict(X_test)
y_pred
Output:
array([0, 0, 0, 0, 0, 0, 1, 0, 1, 0, 0, 0, 1, 0, 1, 0, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1, 0, 1,
           1, 1, 0, 1, 0, 0, 0, 0, 0, 1, 1, 0, 0, 0, 0, 0, 0, 1, 0, 1, 0, 0,
          1, 0, 1, 1, 0, 1, 1, 0, 0, 1, 1, 1, 0, 0, 1, 0, 0, 1, 0, 1, 0, 1,
          0, 0, 1, 1, 1, 0, 0, 1, 0, 0, 1, 1, 1, 1, 1, 1, 0, 0, 1, 1, 1, 1,
          1, 1, 1, 1, 0, 0, 0, 0, 0, 1, 1, 1], dtype=int64)
from sklearn.metrics import confusion_matrix
cm = confusion matrix(y test, y pred)
cm
Output:
array([[53, 15],
                       [ 1, 31]], dtype=int64)
from sklearn.metrics import accuracy score
print("The accuracy score is:", accuracy score(y pred, y test))
```

The accuracy score is: 0.84

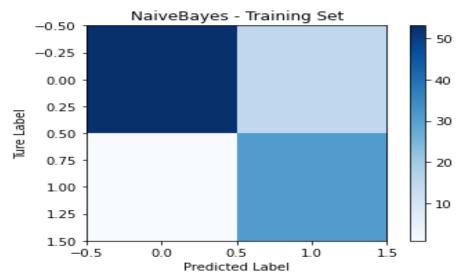
from sklearn.metrics import classification\_report print("classification\_report:") print( classification\_report(y\_pred, y\_test))

### Output:

classification\_report:

	precision	recall	f1-score	support
0	0.78	0.98	0.87	54
1	0.97	0.67	0.79	46
accuracy			0.84	100
macro avg	0.87	0.83	0.83	100
weighted avg	0.87	0.84	0.83	100

plt.imshow(cm,interpolation='nearest',cmap=plt.cm.Blues)
plt.title('NaiveBayes - Training Set')
plt.xlabel('Predicted Label')
plt.ylabel('Ture Label')
plt.colorbar()
plt.show()



# 9. Write a program to implement a Support Vector Machine (SVM) classifier sample training dataset. Fine-tune various hyperparameters and assess the classifier's performance on the dataset.

import numpy as np import matplotlib.pyplot as plt import pandas as pd from sklearn.model\_selection import train\_test\_split from sklearn.preprocessing import StandardScaler

dataset = pd.read\_csv('Social\_Network\_Ads.csv')
Dataset

### **Output:**

User ID	Gender	Age	EstimatedSalary	Purchased	
0	15624510	Male	19	19000	0
1	15810944	Male	35	20000	0
2	15668575	Female	26	43000	0
3	15603246	Female	27	57000	0
4	15804002	Male	19	76000	0
395	15691863	Female	46	41000	1
396	15706071	Male	51	23000	1
397	15654296	Female	50	20000	1
398	15755018	Male	36	33000	0
399	15594041	Female	49	36000	1

# converting gender column to numeric from sklearn.preprocessing import LabelEncoder label\_encoder=LabelEncoder() dataset['Gender']=label\_encoder.fit\_transform(dataset['Gender']) dataset['Gender'].unique()

### Output:

array([1, 0])

# to include gender
X = dataset.iloc[:, [1, 3]].values

```
y = dataset.iloc[:,4].values
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.25, random_state=0)
# X = dataset.iloc[:, [2, 3]].values
# y = dataset.iloc[:,4].values
# X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.25, random_state=0)
# sc = StandardScaler()
# X train = sc.fit transform(X train)
# X test = sc.transform(X test)
from sklearn.preprocessing import MinMaxScaler
scaler = MinMaxScaler()
X train = scaler.fit transform(X train)
X test = scaler.transform(X test)
from sklearn.svm import SVC
classifier = SVC(kernel='poly', random_state=0)
classifier.fit(X train, y train)
Output:
SVC(kernel='poly', random_state=0)
y_pred = classifier.predict(X_test)
y_pred
Output:
array([0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 1, 0, 0, 1, 0, 0, 1,
   1, 0, 0, 0, 0, 1, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
   0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 1, 1, 0, 0, 0, 0, 0, 1,
   0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 1], dtype=int64)
from sklearn.metrics import confusion matrix
```

cm = confusion\_matrix(y\_test, y\_pred)

cm

### Output:

```
array([[66, 2], [19, 13]], dtype=int64)
```

from sklearn.metrics import accuracy\_score
print("The accuracy score is:", accuracy\_score(y\_pred, y\_test))

### Output:

The accuracy score is: 0.79

from sklearn.metrics import classification\_report
print("classification\_report:")
print( classification\_report(y\_pred, y\_test))

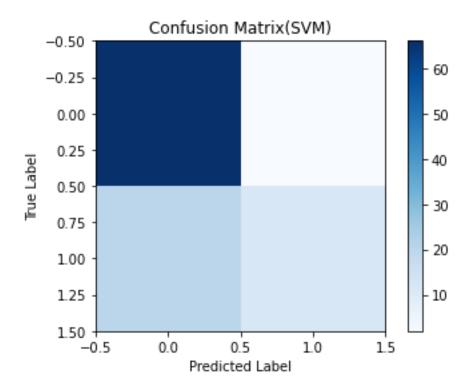
### Output:

classification\_report:

	precision	recall	f1-score	support
0	0.97	0.78	0.86	85
1	0.41	0.87	0.55	15
accuracy			0.79	100
macro avg	0.69	0.82	0.71	100
weighted avg	0.89	0.79	0.82	100

```
plt.imshow(cm,interpolation='nearest',cmap=plt.cm.Blues)
plt.title('Confusion Matrix(SVM)')
plt.xlabel('Predicted Label')
plt.ylabel('True Label')
plt.colorbar()
```

# Show the plot plt.show()



### 10. Write a program to implement Decision Tree classifier. Experiment with different hyperparameters to evaluate and optimize the classifier's performance.

```
import numpy as np
import matplotlib.pyplot as plt
import pandas as pd
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import StandardScaler
```

dataset = pd.read\_csv('Social\_Network\_Ads.csv') X = dataset.iloc[:, [2, 3]].values

y = dataset.iloc[:,4].values

X\_train, X\_test, y\_train, y\_test = train\_test\_split(X, y, test\_size=0.25, random\_state=0)

sc = StandardScaler()

X train = sc.fit transform(X train)

X\_test = sc.transform(X\_test)

dataset

	UserID	Gender	Age	EstimatedSalary	Purchased
0	15624510	Male	19	19000	0
U	13024310	iviale	19	19000	U
1	15810944	Male	35	20000	0
2	15668575	Female	26	43000	0
3	15603246	Female	27	57000	0
4	15804002	Male	19	76000	0
395	15691863	Female	46	41000	1
396	15706071	Male	51	23000	1
397	15654296	Female	50	20000	1
398	15755018	Male	36	33000	0
399	15594041	Female	49	36000	1

```
from sklearn.tree import DecisionTreeClassifier
classifier = DecisionTreeClassifier(criterion='gini', random_state=0)
classifier.fit(X_train, y_train)
Output:
DecisionTreeClassifier(random_state=0)
y pred = classifier.predict(X test)
y_pred
Output:
array([0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 1, 0, 1, 1, 0, 1, 0, 0, 0,
    0, 1, 0, 1, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0,
    1, 0, 0, 0, 0, 1, 1, 0, 0, 1, 1, 1, 0, 0, 1, 0, 0, 1, 0, 1, 0, 1,
    0, 0, 0, 1, 1, 0, 0, 1, 0, 0, 0, 0, 1, 1, 1, 1, 1, 0, 0, 1, 0, 0, 1,
    1, 0, 0, 1, 0, 0, 0, 1, 0, 1, 1, 1], dtype=int64)
from sklearn.metrics import confusion matrix
cm = confusion matrix(y test, y pred)
cm
Output:
array([[62, 6],
    [4, 28]], dtype=int64)
```

plot\_tree(classifier, feature\_names=['Age', 'EstimatedSalary'], class\_names=['0', '1'],

from sklearn.tree import plot\_tree

plt.figure(figsize=(50,50))

filled=True) plt.show()

# 11. Write a program to implement K-means clustering. Using data visualization (Scatter Plot) technique to illustrate the clustering.

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
dataset = pd.read_csv('Mall_Customers.csv')
dataset
```

### **Output:**

	CustomerID	Genre	Age	Annual Income (k\$)	Spending Score (1-100)
0	1	Male	19	15	39
1	2	Male	21	15	81
2	3	Female	20	16	6
3	4	Female	23	16	77
4	5	Female	31	17	40
195	196	Female	35	120	79
196	197	Female	45	126	28
197	198	Male	32	126	74
198	199	Male	32	137	18
199	200	Male	30	137	83

200 rows × 5 columns

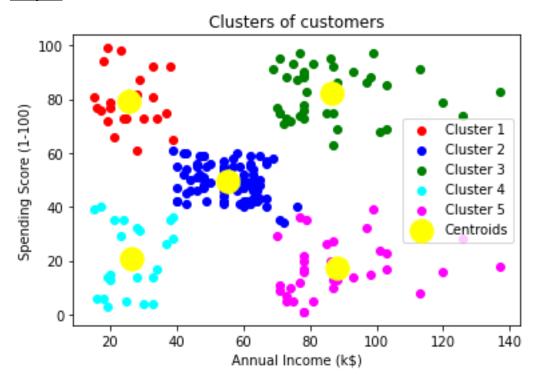
X = dataset.iloc[:, [3, 4]].values

```
# Fitting K-Means to the dataset
from sklearn.cluster import KMeans
kmeans = KMeans(n_clusters=5, init='k-means++', random_state=42)
y_kmeans = kmeans.fit_predict(X)
```

```
# Visualising the clusters
```

```
plt.scatter(X[y_kmeans == 0, 0], X[y_kmeans == 0, 1], c = 'red', label = 'Cluster 1')
plt.scatter(X[y_kmeans == 1, 0], X[y_kmeans == 1, 1], c = 'blue', label = 'Cluster 2')
plt.scatter(X[y_kmeans == 2, 0], X[y_kmeans == 2, 1], c = 'green', label = 'Cluster 3')
plt.scatter(X[y_kmeans == 3, 0], X[y_kmeans == 3, 1], c = 'cyan', label = 'Cluster 4')
plt.scatter(X[y_kmeans == 4, 0], X[y_kmeans == 4, 1], c = 'magenta', label = 'Cluster 5')
plt.scatter(kmeans.cluster_centers_[:, 0], kmeans.cluster_centers_[:, 1], s = 300, c = 'yellow', label = 'Centroids')
```

```
plt.title('Clusters of customers')
plt.xlabel('Annual Income (k$)')
plt.ylabel('Spending Score (1-100)')
plt.legend()
plt.show()
```



# 12. Write a program to implement hierarchical clustering algorithm. Using data visualization (Scatter Plot) technique to illustrate the clustering.

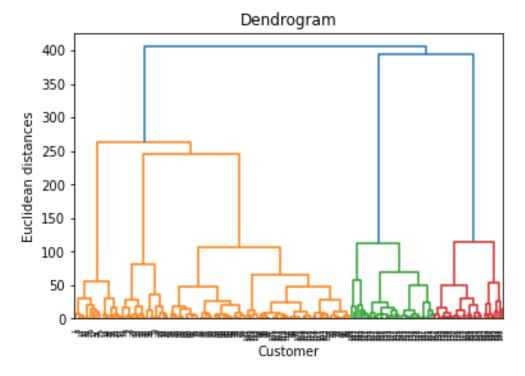
import numpy as np
import matplotlib.pyplot as plt
import pandas as pd
dataset = pd.read\_csv("Mall\_Customers.csv")
dataset.head()

### Output:

	CustomerI D	Genre	Age	Annual Income (k\$)	Spending Score (1- 100)
0	1	Male	19	15	39
1	2	Male	21	15	81
2	3	Female	20	16	6
3	4	Female	23	16	77
4	5	Female	31	17	40

x = dataset.iloc[:,[3,4]].values

import scipy.cluster.hierarchy as sch
dendrogram = sch.dendrogram(sch.linkage(x, method = 'ward'))
plt.title("Dendrogram")
plt.xlabel("Customer")
plt.ylabel("Euclidean distances")
plt.show()



```
from sklearn.cluster import AgglomerativeClustering hc = AgglomerativeClustering(n_clusters=5, linkage = 'ward')  
y_hc = hc.fit_predict(x)  
plt.scatter(x[y_hc == 0, 0], x[y_hc == 0, 1], s = 100, c = "red", label = "cluser 1")  
plt.scatter(x[y_hc == 1, 0], x[y_hc == 1, 1], s = 100, c = "blue", label = "cluser 2")  
plt.scatter(x[y_hc == 2, 0], x[y_hc == 2, 1], s = 100, c = "green", label = "cluser 3")  
plt.scatter(x[y_hc == 3, 0], x[y_hc == 3, 1], s = 100, c = "cyan", label = "cluser 4")  
plt.scatter(x[y_hc == 4, 0], x[y_hc == 4, 1], s = 100, c = "orange", label = "cluser 5")  
plt.title("Clusters of customers")  
plt.title("Annual Income")  
plt.ylabel("Spending Score(1-100)")  
plt.legend()  
plt.show()
```



# 13. Implement density-based clustering using a suitable dataset. Explore the DBSCAN algorithm and visualize the data.

import pandas as pd from sklearn.cluster import DBSCAN import matplotlib.pyplot as plt from sklearn.datasets import make\_blobs

data = pd.read\_csv('blobs.csv')
data

### Output:

	0	1
0	8.622185	1.935796
1	-4.736710	-7.970958
2	9.621222	0.925423
3	6.162095	-0.273254
4	8.697488	-1.057452
0	8.622185	1.935796
•••	•••	•••
995	8.993880	2.203768
996	-5.082768	-9.644539
997	-6.252268	-8.412482
998	-5.479154	-10.536955
999	6.120559	0.968963

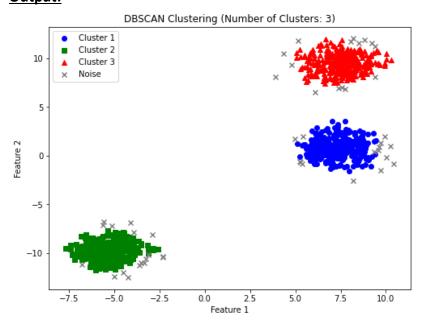
# Extract the features (assuming your CSV file has columns 'Feature1' and 'Feature2')

X = data.iloc[:,[0,1]].values

X

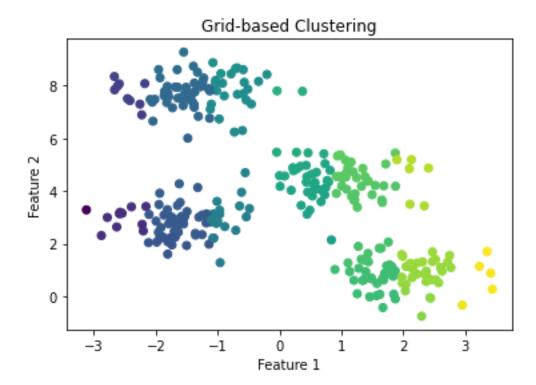
```
array([[ 8.62218539, 1.93579579],
       [ -4.73670958, -7.97095765],
       [ 9.62122205, 0.92542315],
       ...,
       [ -6.2522678, -8.412482 ],
```

```
[-5.479154,-10.53695547],
    [ 6.12055883, 0.96896287]])
# DBSCAN clustering
db = DBSCAN(eps=0.5, min_samples=5)
y_db = db.fit_predict(X)
# Number of clusters in labels, ignoring noise if present (-1)
n clusters = len(set(y db)) - (1 if -1 in y db else 0)
# Plot the clusters
plt.figure(figsize=(8, 6))
plt.scatter(X[y db == 0][:, 0], X[y db == 0][:, 1], c='blue', marker='o', label='Cluster 1')
plt.scatter(X[y_db == 1][:, 0], X[y_db == 1][:, 1], c='green', marker='s', label='Cluster 2')
plt.scatter(X[y db == 2][:, 0], X[y db == 2][:, 1], c='red', marker='^', label='Cluster 3')
plt.scatter(X[y_db == -1][:, 0], X[y_db == -1][:, 1], c='gray', marker='x', label='Noise')
plt.legend(loc='best')
plt.title(f"DBSCAN Clustering (Number of Clusters: {n_clusters_})")
plt.xlabel("Feature 1")
plt.ylabel("Feature 2")
plt.show()
```



# 14. Write a program to implement grid-based clustering using a suitable dataset. Visualize the data using scatter plot.

```
import numpy as np
import matplotlib.pyplot as plt
from sklearn.datasets import make blobs
# Generate synthetic data
data, = make blobs(n samples=300, centers=4, cluster std=.60, random state=0)
# Set the grid size (you can adjust this based on your data distribution)
grid size = 1.0
# Get the minimum and maximum values for x and y coordinates
x min, x max = data[:, 0].min(), data[:, 0].max()
y min, y max = data[:, 1].min(), data[:, 1].max()
# Create a grid by defining intervals using the minimum and maximum values
x \text{ grid} = \text{np.arange}(x \text{ min, } x \text{ max} + \text{grid size, } \text{grid size})
y grid = np.arange(y min, y max + grid size, grid size)
# Initialize labels array with zeros
labels = np.zeros(data.shape[0], dtype=int)
# Assign each data point to a grid cell based on its coordinates
for i, point in enumerate(data):
  x, y = point
  x label = np.searchsorted(x grid, x) - 1
  y_label = np.searchsorted(y_grid, y) - 1
  labels[i] = x_label * len(y_grid) + y_label
# Visualize the clusters
plt.scatter(data[:, 0], data[:, 1], c=labels, cmap='viridis')
plt.title('Grid-based Clustering')
plt.xlabel('Feature 1')
plt.ylabel('Feature 2')
plt.show()
```



### 15. Write a program to perform linear regression using

### i. Single variable

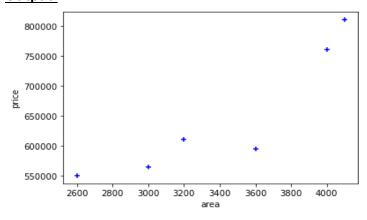
### Ii. Multiple variable

#Using single variable
import pandas as pd
from sklearn import linear\_model
import matplotlib.pyplot as plt
df = pd.read\_csv("homeprices.csv")
df

### **Output:**

	Area	price
0	2600	550000
1	3000	565000
2	3200	610000
3	3600	595000
4	4000	760000
5	4100	810000

plt.xlabel('area')
plt.ylabel('price')
plt.scatter(df.area, df.price, color = 'blue', marker = '+')



```
reg = linear_model.LinearRegression()
reg.fit(df[['area']], df.price)
reg.coef_
```

array([167.30954677])

reg.intercept\_

### Output:

76692.3818707813

reg.predict([[6800]])

### Output:

array([1214397.29990357])

#Using multiple variable

import pandas as pd
from sklearn import linear\_model
import matplotlib.pyplot as plt
df = pd.read\_csv("homeprices\_multiple.csv")
df

### Output:

	area	bedrooms	age	price
0	2600	3.0	20	550000
1	3000	4.0	15	565000
2	3200	NaN	18	610000
3	3600	3.0	30	595000
4	4000	5.0	8	760000
5	4100	6.0	8	810000

df.bedrooms.median()

### **Output:**

4.0

```
df.bedrooms = df.bedrooms.fillna(df.bedrooms.median())
df
```

	area	bedrooms	age	price
0	2600	3.0	20	550000
1	3000	4.0	15	565000
2	3200	4.0	18	610000
3	3600	3.0	30	595000
4	4000	5.0	8	760000
5	4100	6.0	8	810000

```
x = df.iloc[:, [0,1,2]].values
y = df.iloc[:, 3].values
reg = linear_model.LinearRegression()
reg.fit(x, y)
```

### **Output:**

LinearRegression()

reg.coef\_

### Output:

array([ 112.06244194, 23388.88007794, -3231.71790863])

reg.intercept\_

### Output:

221323.00186540408

reg.predict([[2600, 3, 30]])

### Output:

array([485900.45388978])

# 16. Write a program to implement chi-square test for feature selection to train SVM classifier using suitable dataset.

```
import pandas as pd
# Load dataset
data = pd.read csv('fruit data with colours.csv')
data.head(5)
fruit label = 'fruit label'
fruit subtype = 'fruit subtype'
fruit name = 'fruit name'
# Drop non-numeric columns if necessary and extract features (X) and target (y)
X = data.drop([fruit label, fruit subtype, fruit name], axis=1) # Features
y = data[fruit label]
from sklearn.feature selection import SelectKBest, chi2
k selected features = 4 # Adjust this value based on how many top features you want to select
chi2 selector = SelectKBest(chi2, k=k selected features)
X selected = chi2 selector.fit transform(X, y)
from sklearn.model selection import train test split
X train, X test, y train, y test = train test split(X selected, y, test size=0.2, random state=42)
# Step 3: Train the SVM classifier
from sklearn.svm import SVC
svm classifier = SVC(kernel='linear')
svm_classifier.fit(X_train, y_train)
SVC(kernel='linear')
# Step 4: Evaluate the SVM classifier
from sklearn.metrics import accuracy score, classification report
y pred = svm classifier.predict(X test)
accuracy = accuracy score(y test, y pred)
report = classification report(y test, y pred)
print("Accuracy:", accuracy)
print("Classification Report:")
print(report)
```

Accuracy: 0.75

Classification Report:

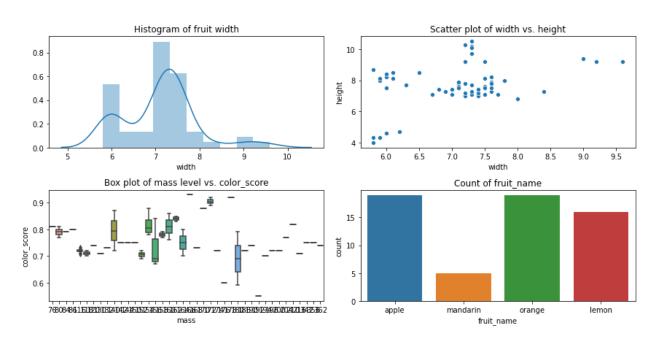
	precision	recall	f1-score	support
1	0.67	0.67	0.67	3
2	1.00	1.00	1.00	2
3	0.33	0.50	0.40	2
4	1.00	0.80	0.89	5
accuracy			0.75	12
macro avg	0.75	0.74	0.74	12
weighted avg	0.81	0.75	0.77	12

## 17. Implement a program to perform various data visualization techniques on sample dataset.

```
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
# Load dataset (replace with your dataset loading code)
data = pd.read_csv('fruit_data_with_colours.csv')
# Display the first few rows of the dataset
print("First few rows of the dataset:")
print(data.head())
plt.figure(figsize=(12, 6))
# Plot 1: Histogram
plt.subplot(2, 2, 1)
sns.distplot(data['width'], bins=10, kde=True)
plt.title('Histogram of fruit width')
# Plot 2: Scatter plot
plt.subplot(2, 2, 2)
sns.scatterplot(x='width', y='height', data=data)
plt.title('Scatter plot of width vs. height')
# Plot 3: Box plot
plt.subplot(2, 2, 3)
sns.boxplot(x='mass', y='color score', data=data)
plt.title('Box plot of mass level vs. color score')
# Plot 4: Count plot
plt.subplot(2, 2, 4)
sns.countplot(x='fruit name', data=data)
plt.title('Count of fruit name')
# Adjust layout
plt.tight_layout()
```

### First few rows of the dataset:

	<pre>fruit_label</pre>	<pre>fruit_name</pre>	<pre>fruit_subtype</pre>	mass	width	height	color_score
0	1	apple	granny_smith	192	8.4	7.3	0.55
1	1	apple	granny_smith	180	8.0	6.8	0.59
2	1	apple	granny_smith	176	7.4	7.2	0.60
3	2	mandarin	mandarin	86	6.2	4.7	0.80
4	2	mandarin	mandarin	84	6.0	4.6	0.79



### 18. Implement a program to perform attribute selection measures.

```
import pandas as pd
import numpy as np
def entropy(data):
  values, counts = np.unique(data, return_counts=True)
  probs = counts / len(data)
  return -np.sum(probs * np.log2(probs))
def information gain(data, attribute index):
  total entropy = entropy(data[:, -1])
  values, counts = np.unique(data[:, attribute index], return counts=True)
  weighted_entropy = sum((counts[i] / len(data)) * entropy(data[data[:, attribute index] ==
values[i], -1]) for i in range(len(values)))
  return total entropy - weighted entropy
def gain_ratio(data, attribute_index):
  # Information Gain
  ig = information gain(data, attribute index)
  # Calculate Intrinsic Value
  values, counts = np.unique(data[:, attribute index], return counts=True)
  total instances = len(data)
  intrinsic value = -np.sum((counts / total_instances) * np.log2(counts / total_instances))
  return ig / intrinsic value if intrinsic value != 0 else 0
def gini index(data):
  # Calculate the Gini index of a dataset
  class labels = data[:, -1]
  total instances = len(class labels)
  label counts = np.unique(class labels, return counts=True)[1]
  label probabilities = label counts / total instances
  gini = 1 - np.sum(label probabilities**2)
  return gini
def gini_index_attribute(data, attribute_index):
  # Calculate the Gini index of an attribute in a dataset
  attribute_values = np.unique(data[:, attribute_index])
```

```
total_instances = len(data)
  gini attribute = 0
  for value in attribute_values:
    subset = data[data[:, attribute_index] == value]
    subset_instances = len(subset)
    gini subset = gini index(subset)
    gini attribute += (subset instances / total instances) * gini subset
  return gini attribute
# Load CSV file
df = pd.read_csv("Buys_Computer.csv")
data = df.values
print("Dataset loaded successfully:")
print(df)
while True:
  print("\n1. Information Gain\n2. Gain Ratio\n3. Gini Index\n4. Exit")
  ch = input("Enter your choice: ")
  if ch == "1":
    try:
      attribute index = int(input(f"Enter the index of the attribute (0 to {data.shape[1] - 2}) for
which you want to calculate Information Gain: "))
      if 0 <= attribute_index < data.shape[1] - 1:
         ig = information_gain(data, attribute_index)
         print(f"Information Gain for attribute {attribute index}: {ig}")
      else:
         print(f"Invalid attribute index. Please enter a number between 0 and {data.shape[1] -
2}.")
    except ValueError:
      print("Invalid input. Please enter a valid integer for the attribute index.")
    except Exception as e:
      print(f"An error occurred: {e}")
  elif ch == "2":
    try:
```

```
attribute index = int(input(f"Enter the index of the attribute (0 to {data.shape[1] - 2}) for
which you want to calculate Gain Ratio: "))
       if 0 <= attribute_index < data.shape[1] - 1:
         # Calculating Gain Ratio for the specified attribute
         gain_ratio_attr = gain_ratio(data, attribute_index)
         print(f"Gain Ratio for attribute {attribute index}: {gain ratio attr}")
         print(f"Invalid attribute index. Please enter a number between 0 and {data.shape[1] -
2}.")
    except ValueError:
       print("Invalid input. Please enter a valid integer for the attribute index.")
    except Exception as e:
       print(f"An error occurred: {e}")
  elif ch == "3":
    try:
       attribute index = int(input(f"Enter the index of the attribute (0 to {data.shape[1] - 2}) for
which you want to calculate Gini Index: "))
       if 0 <= attribute index < data.shape[1] - 1:
         # Calculating Gini index for the specified attribute
         gini_attr = gini_index_attribute(data, attribute_index)
         print(f"Gini index for attribute {attribute index}: {gini attr}")
       else:
         print(f"Invalid attribute index. Please enter a number between 0 and {data.shape[1] -
2}.")
    except ValueError:
       print("Invalid input. Please enter a valid integer for the attribute index.")
    except Exception as e:
       print(f"An error occurred: {e}")
  elif ch == "4":
    break
  else:
    print("Invalid choice")
```

**Dataset loaded successfully:** 

	age	income	student	credit	_rating buys_computer	
0		<=30	high	no	fair	no
1		<=30	high	no	excellent	no
2	31.	40	high	no	fair	yes
3		>40 n	nedium	no	fair	yes
4		>40	low	yes	fair	yes
5		>40	low	yes	excellent	no
6	31.	40	low	yes	excellent	yes
7		<=30 n	nedium	no	fair	no
8		<=30	low	yes	fair	yes
9		>40 n	nedium	yes	fair	yes
10		<=30 n	nedium	yes	excellent	yes
11	31.	40 n	nedium	no	excellent	yes
12	31.	40	high	yes	fair	yes
13		>40 n	nedium	no	excellent	no

- 1. Information Gain
- 2. Gain Ratio
- 3. Gini Index
- 4. Exit

**Enter your choice: 1** 

Enter the index of the attribute (0 to 3) for which you want to calculate Information Gain: 1 Information Gain for attribute 1: 0.02922256565895487

- 1. Information Gain
- 2. Gain Ratio
- 3. Gini Index
- 4. Exit

**Enter your choice: 1** 

Enter the index of the attribute (0 to 3) for which you want to calculate Information Gain: 3 Information Gain for attribute 3: 0.04812703040826949

- 1. Information Gain
- 2. Gain Ratio
- 3. Gini Index
- 4. Exit

Enter your choice: 3

Enter the index of the attribute (0 to 3) for which you want to calculate Gini Index: 0

### Gini index for attribute 0: 0.34285714285714286

- 1. Information Gain
- 2. Gain Ratio
- 3. Gini Index
- 4. Exit

Enter your choice: 2

Enter the index of the attribute (0 to 3) for which you want to calculate Gain Ratio: 2 Gain Ratio for attribute 2: 0.15183550136234159

- 1. Information Gain
- 2. Gain Ratio
- 3. Gini Index
- 4. Exit

Enter your choice: 4

### 19. Implement a program to perform different distance measures.

```
import pandas as pd
import numpy as np
# Function to calculate Euclidean distance
def euclidean distance(instance1, instance2):
  return np.linalg.norm(instance1 - instance2)
# Function to calculate Manhattan distance
def manhattan_distance(instance1, instance2):
  return np.sum(np.abs(instance1 - instance2))
# Function to calculate Cosine similarity
def cosine similarity(instance1, instance2):
  dot product = np.dot(instance1, instance2)
  norm1 = np.linalg.norm(instance1)
  norm2 = np.linalg.norm(instance2)
  return dot product / (norm1 * norm2)
# Load CSV file
file path = 'homeprices multiple.csv' #input("Enter the path to the CSV file: ")
df = pd.read csv(file path)
# Print loaded dataset
print("Dataset loaded successfully:")
print(df)
# Mapping of distance measure names to functions
distance measures = {
  "1": ("Euclidean", euclidean_distance),
  "2": ("Manhattan", manhattan distance),
  "3": ("Cosine Similarity", cosine similarity)
}
# Print distance measure options
print("\nSelect a distance measure:")
for key, (measure_name, _) in distance_measures.items():
```

```
print(f"{key}. {measure_name}")
# Accept user input for selecting distance measure
selected_measure_name = input("Enter the index or name of the distance measure: ")
# Validate the selected measure
if selected measure name in distance measures:
  selected_measure = distance_measures[selected_measure name][1] # Get the function
corresponding to the selected measure
  selected measure name = distance measures[selected measure name][0] # Get the name
of the selected measure
else:
  print("Invalid distance measure selection. Please choose from the available options.")
  exit()
# Input indices of two instances
index1 = int(input(f"Enter index of the first instance (0 to {len(df)-1}): "))
index2 = int(input(f"Enter index of the second instance (0 to {len(df)-1}): "))
# Validate indices
if 0 \le index 1 \le len(df) and 0 \le index 2 \le len(df):
  instance1 = df.iloc[index1, :-1].values # Exclude last column (assuming it's the target variable)
  instance2 = df.iloc[index2, :-1].values # Exclude last column (assuming it's the target variable)
  # Calculate distance based on user's choice
  distance = selected measure(instance1, instance2)
  print(f"{selected measure name} distance between instance {index1} and instance {index2}:
{distance}")
else:
  print(f"Invalid indices. Please enter indices between 0 and {len(df)-1}.")
```

### Output 1:

**Dataset loaded successfully:** 

area bedrooms age price

- 0 2600 3.0 20 550000
- 1 3000 4.0 15 565000
- 2 3200 NaN 18 610000
- 3 3600 3.0 30 595000
- 4 4000 5.0 8 760000
- 5 4100 6.0 8 810000

### Select a distance measure:

- 1. Euclidean
- 2. Manhattan
- 3. Cosine Similarity

Enter the index or name of the distance measure: 1

Enter index of the first instance (0 to 5): 5

Enter index of the second instance (0 to 5): 0

Euclidean distance between instance 5 and instance 0: 1500.0509991330296

### Output 2:

**Dataset loaded successfully:** 

area bedrooms age price

- 0 2600 3.0 20 550000
- 1 3000 4.0 15 565000
- 2 3200 NaN 18 610000
- 3 3600 3.0 30 595000
- 4 4000 5.0 8 760000
- 5 4100 6.0 8 810000

### Select a distance measure:

- 1. Euclidean
- 2. Manhattan
- 3. Cosine Similarity

Enter the index or name of the distance measure: 2

Enter index of the first instance (0 to 5): 0

Enter index of the second instance (0 to 5): 1

Manhattan distance between instance 0 and instance 1: 406.0

# 20. Implement a LinearSVC classifier for text classification using TF-IDF features from char n-grams. Evaluate the performance of the model.

### #With built-in

import pandas as pd
import spacy
import string
import contractions
import re
import nltk
import emoji
from nltk import word\_tokenize
from nltk.corpus import stopwords
from nltk.stem import WordNetLemmatizer
from nltk.stem import PorterStemmer
from num2words import num2words

nltk.download('punkt')
nltk.download('stopwords')
nltk.download('whitespace')
nltk.download('wordnet')
nltk.download('omw-1.4')
stop\_words = set(stopwords.words("english"))
data = pd.read\_csv('Training.tsv',sep='\t')
data.head(10)

	tweet_id	text	label
0	1382343793341575169	@IrvineWelsh I don't know about you Irvine	0
		but	
1	1377631738692796417	I bet money if i went n took a covid test	0
		righ	

0	@JamesMelville My wife received a POSITIVE Cov	1386448010029240326	2
0	Out of the 180,000+ people who have had the tw	1361342676340211717	3
0	My whole family is sick af and here I am now i	1386757983254765569	4
0	@renfrew1962 @PeakePolly @J_Deliciouso I'm not	1382001700853125122	5
1	Test came back positive, no surprise. I have c	1383272654212272136	6
0	My Pawpaw has been in the hospital a few days	1374479299047084035	7
0	@MattHancock 4 people I know had covid and rec	1354020426620547072	8
1	I'm going to sound like I have lost my marbles	1362671045136809985	9

data['label'].value\_counts()

```
Output:
```

0 6266

1 1334

Name: label, dtype: int64

```
ps =PorterStemmer()
lemmatiser = WordNetLemmatizer()
english_stopwords = stopwords.words('english')
exclude = set(string.punctuation)
def preprocess(text):
    #text=demoji.findall(df['Text'])
    text = contractions.fix(text.lower(), slang=True)
    text = re.sub(r'\d+', lambda x: num2words(int(x.group(0))), text)
    #text= re.sub(r'\d+', ", text)
    text=re.sub(r'\s', ", text)
    text=re.sub(r'',", text)
    text=re.sub('<.*?>',",text)
    text=re.sub(r'http\S+', ", text)
    #text=emoji.demojize(text, delimiters=(" ", " "))
```

```
text = ".join(ch for ch in text if ch not in exclude)
  tokens = word_tokenize(text)
  #print("Tokens:", tokens)
  text = [t for t in tokens if t not in english_stopwords]
  text = " ".join(text)
  return text
import emoji
#import demoji
#demoji.download_codes()
def emo(text):
  temp=emoji.demojize(text,delimiters=(" "," "))
  temp=temp.replace("_"," ")
  return temp
data['emo']=data["text"].apply(lambda x:emo(x))
data["clean_text"]=data['emo'].apply(lambda X: preprocess(X))
data.head()
```

	tweet_id	text	label	emo	clean_text
0	13823437933	@IrvineWelsh I don't	0	@IrvineWelsh I	irvinewelsh know
	41575169	know about you		don't know about	irvine keep told
		Irvine but		you Irvine but	covid exist
1	13776317386	I bet money if i went	0	I bet money if i	bet money went n
	92796417	n took a covid test		went n took a	took covid test right
		righ		covid test righ	going t
2	13864480100	@JamesMelville My	0	@JamesMelville	jamesmelville wife
	29240326	wife received a		My wife received	received positive
		POSITIVE Cov		a POSITIVE Cov	covid tes
3	13613426763	Out of the 180,000+	0	Out of the	one hundred
	40211717	people who have had		180,000+ people	eightyzero people
		the tw		who have had the	two vaccine shot
				tw	
4	13867579832	My whole family is	0	My whole family	whole family sick af
	54765569	sick af and here I am		is sick af and here	hospital heart
		now i		I am now i	palpitatio

from sklearn.model selection import train test split

X\_train, X\_test, y\_train, y\_test = train\_test\_split(data['clean\_text'], data['label'], test\_size=0.33, random state=42)

### Feature Extraction: TF-IDF (char\_wb)

```
Tfidf_vec1 = TfidfVectorizer(analyzer='char_wb', ngram_range=(1, 5), max_df=1.0, min_df=1, max_features=5000) count_train1 = Tfidf_vec1.fit(X_train) train_features1 = Tfidf_vec1.transform(X_train) test features1 = Tfidf_vec1.transform(X_test)
```

### Feature Extraction- (word) TFIDF

```
Tfidf_vec2 = TfidfVectorizer(analyzer='word', ngram_range=(1, 3), max_df=1.0, min_df=1, max_features=5000)

count_train2 = Tfidf_vec2.fit(X_train)

train_features2 = Tfidf_vec2.transform(X_train)

test_features2 = Tfidf_vec2.transform(X_test)
```

### Feature Extraction - (word) CountVectorizer

```
count_vec3 = CountVectorizer(analyzer='word', ngram_range=(1, 3), max_df=1.0, min_df=1,
max_features=5000)
count_train3 = count_vec3.fit(X_train)
train_features3 = count_vec3.transform(X_train)
test_features3 = count_vec3.transform(X_test)
```

### Feature Extraction - (char\_wb) CountVectorizer

```
count_vec4 = CountVectorizer(analyzer='char_wb', ngram_range=(1, 5), max_df=1.0, min_df=1,
max_features=5000)
count_train4 = count_vec4.fit(X_train)
train_features4 = count_vec4.transform(X_train)
test_features4 = count_vec4.transform(X_test)
```

### Model Building with SVM - LinearSVC

clf1 =LinearSVC(C=1.0, class weight="balanced", max iter=10000, random state=123)

```
clf1.fit(train_features1, y_train)
y_pred1=clf1.predict(test_features1)
accuracy = accuracy_score(y_test, y_pred1)
print("Test Accuracy(Feature Extraction: TF-IDF (char_wb)):", round(accuracy*100, 4))
print("\n", classification_report(y_test, y_pred1))
```

Test Accuracy(Feature Extraction: TF-IDF (char\_wb)): 79.1866

	precision	recall	f1-score	support
0	0.92	0.81	0.86	2040
1	0.46	0.70	0.56	468
accuracy			0.79	2508
macro avg	0.69	0.76	0.71	2508
weighted avg	0.84	0.79	0.81	2508

```
clf1 =LinearSVC(C=1.0, class_weight="balanced", max_iter=10000, random_state=123)
clf1.fit(train_features2, y_train)
y_pred2=clf1.predict(test_features2)
accuracy = accuracy_score(y_test, y_pred2)
print("Test Accuracy(Feature Extraction- (word) TFIDF):", round(accuracy*100, 4))
print("\n", classification_report(y_test, y_pred2))
```

### Output:

Test Accuracy(Feature Extraction- (word) TFIDF): 81.2998

	precision	recall	f1-score	support
0	0.91	0.86	0.88	2040
1	0.50	0.62	0.55	468
accuracy			0.81	2508
macro avg	0.70	0.74	0.72	2508
weighted avg	0.83	0.81	0.82	2508

clf1=LinearSVC(C=1.0, class\_weight="balanced", max\_iter=10000,random\_state=123) clf1.fit(train\_features3, y\_train)

```
y_pred3 = clf1.predict(test_features3)
accuracy = accuracy_score(y_test, y_pred3)
print("Test Accuracy(Feature Extraction - (word) CountVectorizer):", round(accuracy*100, 4))
print("\n", classification_report(y_test, y_pred3))
```

Test Accuracy(Feature Extraction - (word) CountVectorizer): 80.2632

	precision	recall	f1-score	support
0	0.89	0.87	0.88	2040
1	0.47	0.51	0.49	468
accuracy			0.80	2508
macro avg	0.68	0.69	0.69	2508
weighted avg	0.81	0.80	0.81	2508

clf1 = LinearSVC(C=1.0, class\_weight="balanced", max\_iter=10000, random\_state=123)
clf1.fit(train\_features4, y\_train)
y\_pred4 = clf1.predict(test\_features4)
accuracy = accuracy\_score(y\_test, y\_pred4)
print("Test AccuracyFeature Extraction - (char\_wb) CountVectorizer:", round(accuracy\*100, 4))
print("\n", classification\_report(y\_test, y\_pred4))

### **Output:**

Test AccuracyFeature Extraction - (char\_wb) CountVectorizer: 79.7448

	precision	recall	f1-score	support
0	0.88	0.86	0.87	2040
1	0.46	0.50	0.48	468
accuracy			0.80	2508
macro avg	0.67	0.68	0.68	2508
weighted avg	0.80	0.80	0.80	2508