

Program 1: Write programs to perform basic array creation, operations, reshaping, indexing, and some statistical operations using NumPy

1) **Basic Array Creation**

```
import numpy as np

# Creating an array from a list
array_from_list = np.array([1, 2, 3, 4, 5])
print("Array from list:", array_from_list)

# Creating an array with a range of values
array_with_range = np.arange(10)
print("Array with range:", array_with_range)

# Creating an array of zeros
zeros_array = np.zeros((3, 3))
print("Array of zeros:\n", zeros_array)

# Creating an array of ones
ones_array = np.ones((2, 5))
print("Array of ones:\n", ones_array)
```

Output

```
Array from list: [1 2 3 4 5]
Array with range: [0 1 2 3 4 5 6 7 8 9]
Array of zeros:
[[0. 0. 0.]
 [0. 0. 0.]
 [0. 0. 0.]]
Array of ones:
[[1. 1. 1. 1. 1.]
 [1. 1. 1. 1. 1.]]
```

2. Array Operations

import numpy as np

Creating arrays

a = np.array([1, 2, 3])

b = np.array([4, 5, 6])

Element-wise addition

add_result = np.add(a, b)

print("Addition:", add_result)

Element-wise subtraction

sub_result = np.subtract(a, b)

print("Subtraction:", sub_result)

Element-wise multiplication

mul_result = np.multiply(a, b)

print("Multiplication:", mul_result)

Element-wise division

div_result = np.divide(a, b)

print("Division:", div_result)

Output

Addition: [5 7 9]

Subtraction: [-3 -3 -3]

Multiplication: [4 10 18]

Division: [0.25 0.4 0.5]

3. Array Reshaping

```
import numpy as np
```

```
# Creating a 1D array  
array = np.arange(12)  
print("Original array:", array)
```

```
# Reshaping to 2D array (3x4)  
reshaped_array = array.reshape((3, 4))  
print("Reshaped to 3x4:\n", reshaped_array)
```

```
# Reshaping to 3D array (2x3x2)  
reshaped_array_3d = array.reshape((2, 3, 2))  
print("Reshaped to 2x3x2:\n", reshaped_array_3d)
```

output

```
Original array: [ 0  1  2  3  4  5  6  7  8  9 10 11]  
Reshaped to 3x4:  
[[ 0  1  2  3]  
 [ 4  5  6  7]  
 [ 8  9 10 11]]  
Reshaped to 2x3x2:  
[[[ 0  1]  
   [ 2  3]  
   [ 4  5]]  
  
 [[ 6  7]  
   [ 8  9]  
  [10 11]]]
```

4.Indexing and Slicing

import numpy as np

Creating a 2D array

array_2d = np.array([[1, 2, 3], [4, 5, 6], [7, 8, 9]])

print("2D Array:\n", array_2d)

Accessing element at (1, 2)

element = array_2d[1, 2]

print("Element at (1, 2):", element)

Slicing a subarray (rows 1 to 2, columns 0 to 1)

subarray = array_2d[1:3, 0:2]

print("Sliced subarray:\n", subarray)

Accessing a column (all rows, column 1)

column = array_2d[:, 1]

print("Column 1:", column)

output

```
2D Array:
[[1 2 3]
 [4 5 6]
 [7 8 9]]
Element at (1, 2): 6
Sliced subarray:
[[4 5]
 [7 8]]
Column 1: [2 5 8]
```

5. Statistical Operations

import numpy as np

Creating an array

array = np.array([1, 2, 3, 4, 5, 6, 7, 8, 9, 10])

Calculating mean

mean = np.mean(array)

print("Mean:", mean)

Calculating median

median = np.median(array)

print("Median:", median)

Calculating standard deviation

std_dev = np.std(array)

print("Standard Deviation:", std_dev)

Finding minimum and maximum values

min_val = np.min(array)

max_val = np.max(array)

print("Minimum value:", min_val)

print("Maximum value:", max_val)

output

Mean: 5.5

Median: 5.5

Standard Deviation: 2.8722813232690143

Minimum value: 1

Maximum value: 10

PROGRAM 2:Create two 2D arrays using array object and

- a. Add the 2 matrices and print it
- b. Subtract 2 matrices
- c. Multiply the individual elements of matrix
- d. Divide the elements of the matrices
- e. Perform matrix multiplication
- f. Display transpose of the matrix
- g. Sum of diagonal elements of a matrix

```
import numpy as np
matrix1 = np.array([[1, 2, 3], [4, 5, 6], [7, 8, 9]])
matrix2 = np.array([[9, 8, 7], [6, 5, 4], [3, 2, 1]])
matrix_sum = matrix1 + matrix2
matrix_diff = matrix1 - matrix2
matrix_product = matrix1 * matrix2
matrix_divide = matrix1 / matrix2
matrix_multiply = np.dot(matrix1, matrix2)
matrix1_transpose = np.transpose(matrix1)
diagonal_sum = np.trace(matrix1)
print("Matrix 1:\n", matrix1)
print("Matrix 2:\n", matrix2)
print("Matrix Sum:\n", matrix_sum)
print("Matrix Difference:\n", matrix_diff)
print("Matrix Element-wise Product:\n", matrix_product)
print("Matrix Element-wise Division:\n", matrix_divide)
print("Matrix Multiplication:\n", matrix_multiply)
print("Transpose of Matrix 1:\n", matrix1_transpose)
print("Sum of Diagonal Elements of Matrix 1:", diagonal_sum)
```

output

```
▶ Matrix 1:  
↔ [[1 2 3]  
   [4 5 6]  
   [7 8 9]]  
Matrix 2:  
   [[9 8 7]  
    [6 5 4]  
    [3 2 1]]  
Matrix Sum:  
   [[10 10 10]  
    [10 10 10]  
    [10 10 10]]  
Matrix Difference:  
   [[-8 -6 -4]  
    [-2  0  2]  
    [ 4  6  8]]  
Matrix Element-wise Product:  
   [[ 9 16 21]  
    [24 25 24]  
    [21 16  9]]  
Matrix Element-wise Division:  
   [[0.11111111 0.25      0.42857143]  
    [0.66666667 1.        1.5       ]  
    [2.33333333 4.        9.        ]]  
Matrix Multiplication:  
   [[ 30  24  18]  
    [ 84  69  54]  
    [138 114  90]]  
Transpose of Matrix 1:  
   [[1 4 7]  
    [2 5 8]  
    [3 6 9]]  
Sum of Diagonal Elements of Matrix 1: 15
```

PROGRAM 3:

Write a program to display the elements of the matrix X to different powers and identitymatrix of a given matrix .Also create another matrix Y with same dimensions and display X^2+2Y

```
import numpy as np;
X = np.array([[1, 2],
              [3, 4]])
Y = np.array([[5,6], [7,8]])
print("Matrix X is :\n ",X)
print("Matrix Y is : \n",Y)
a=np.power(X,2)
print("X^2=",a)
result = a + 2 * Y
print("X^2+2*Y is \n ",result)
```

output

```
⇒ Matrix X is :
  [[1 2]
   [3 4]]
Matrix Y is :
  [[5 6]
   [7 8]]
X^2= [[ 1  4]
      [ 9 16]]
X^2+2*Y is
  [[11 16]
   [23 32]]
```


PROGRAM 4:

Create a 2 Dimensional array with 4 rows and 4 columns.

- a. Display all elements excluding the first row
- b. Display all elements excluding the last column
- c. Display the elements of 1st and 2nd column in 2nd and 3rd row
- d. Display the elements of 2nd and 3rd column
- e. Display 2nd and 3rd element of 1st row
- f. Display the elements from indices 4 to 10 in descending order

```
import numpy as np
two_dimensional_array = np.array([[1, 2, 3, 4],
[5, 6, 7, 8],
[9, 10, 11, 12],
[13, 14, 15, 16]])
excluding_first_row = two_dimensional_array[1:]
excluding_last_column = two_dimensional_array[:, :-1]
column_1_2_in_row_2_3 = two_dimensional_array[1:3, 0:2]
column_2_3 = two_dimensional_array[:, 1:3]
elements_2_3_in_first_row = two_dimensional_array[0, 1:3]
descending_order = two_dimensional_array.ravel()[::-1][4:11]
print("Original 2D array:\n", two_dimensional_array)
print("Elements excluding the first row:\n", excluding_first_row)
print("Elements excluding the last column:\n", excluding_last_column)
print("Elements of the 1st and 2nd column in the 2nd and 3rd row:\n",
column_1_2_in_row_2_3)
print("Elements of the 2nd and 3rd column:\n", column_2_3)
print("2nd and 3rd element of the 1st row:\n", elements_2_3_in_first_row)
print("Elements from indices 4 to 10 in descending order:\n", descending_order)
```

output

```
⇒ Original 2D array:
[[ 1  2  3  4]
 [ 5  6  7  8]
 [ 9 10 11 12]
 [13 14 15 16]]
Elements excluding the first row:
[[ 5  6  7  8]
 [ 9 10 11 12]
 [13 14 15 16]]
Elements excluding the last column:
[[ 1  2  3]
 [ 5  6  7]
 [ 9 10 11]
 [13 14 15]]
Elements of the 1st and 2nd column in the 2nd and 3rd row:
[[ 5  6]
 [ 9 10]]
Elements of the 2nd and 3rd column:
[[ 2  3]
 [ 6  7]
 [10 11]
 [14 15]]
2nd and 3rd element of the 1st row:
[2 3]
Elements from indices 4 to 10 in descending order:
[12 11 10  9  8  7  6]
```

PROGRAM 5:

Given a matrix-vector equation $AX=b$. Write a program to find out the value of X using `solve()`, given A and b as below

$$X=A^{-1} b.$$

$$A = \begin{bmatrix} 2 & 1 & -2 \\ 3 & 0 & 1 \\ 1 & 1 & -1 \end{bmatrix} \quad \mathbf{b} = \begin{bmatrix} -3 \\ 5 \\ -2 \end{bmatrix}$$

`np.linalg.solve(A, b)` is a function in the NumPy library used to solve a system of linear equations of the form $AX = b$, where:

- A is a square matrix (with shape $n \times n$).
- b is a vector or matrix (with shape n or $n \times m$) representing the right-hand side of the equation.

```
import numpy as np
A = np.array([[2, 1,-2],[3,0,1],[1,1,-1]])
b = np.array([-3,5,-2])
X = np.linalg.solve(A, b)
print("Matrix A:")
print(A)
print("Vector b:")
print(b)
print("Solution for X:")
print(X)
```

Output

```
⇒ Matrix A:
[[ 2  1 -2]
 [ 3  0  1]
 [ 1  1 -1]]
Vector b:
[-3  5 -2]
Solution for X:
[ 1. -1.  2.]
```

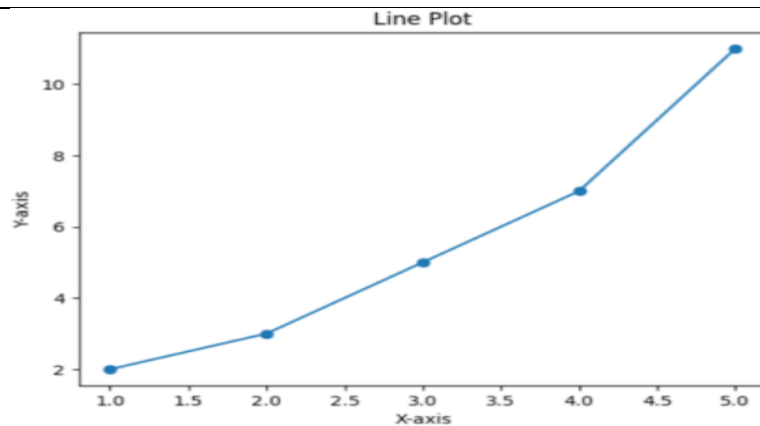
Matplotlib

Matplotlib is a low level graph plotting library in python that serves as a visualization utility.

Program 6: Different Types of Plots using Matplotlib

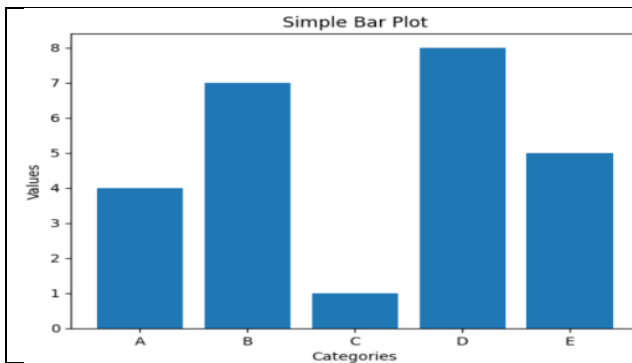
1) Line Plot

```
import matplotlib.pyplot as plt
x = [1, 2, 3, 4, 5]
y = [2, 3, 5, 7, 11]
plt.plot(x, y, marker='o') # Create line plot
plt.title('Line Plot')
plt.xlabel('X-axis')
plt.ylabel('Y-axis')
plt.show()
```

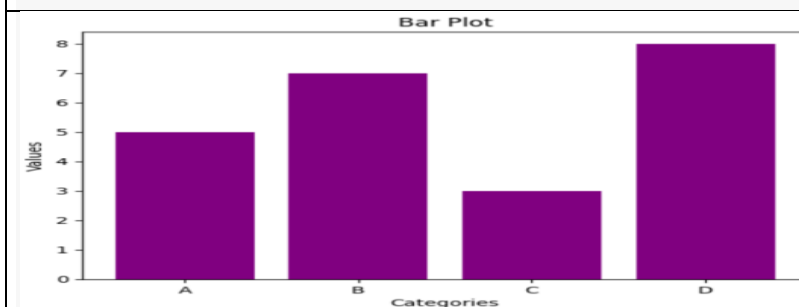


2) Bar Plot

```
import matplotlib.pyplot as plt
categories = ['A', 'B', 'C', 'D', 'E']
values = [4, 7, 1, 8, 5]
plt.bar(categories, values) # Create a bar plot
# Add labels and title
plt.xlabel('Categories')
plt.ylabel('Values')
plt.title('Simple Bar Plot')
plt.show()
```

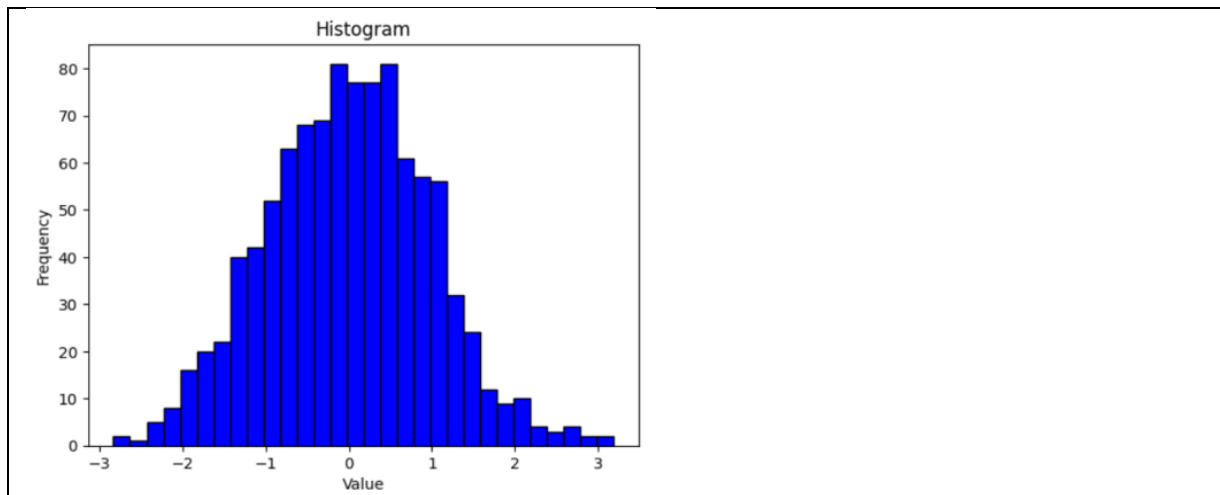


```
# Create bar plot
import matplotlib.pyplot as plt
categories = ['A', 'B', 'C', 'D']
values = [5, 7, 3, 8]
plt.bar(categories, values, color='purple') # Create bar plot
plt.title('Bar Plot')
plt.xlabel('Categories')
plt.ylabel('Values')
plt.show()
```



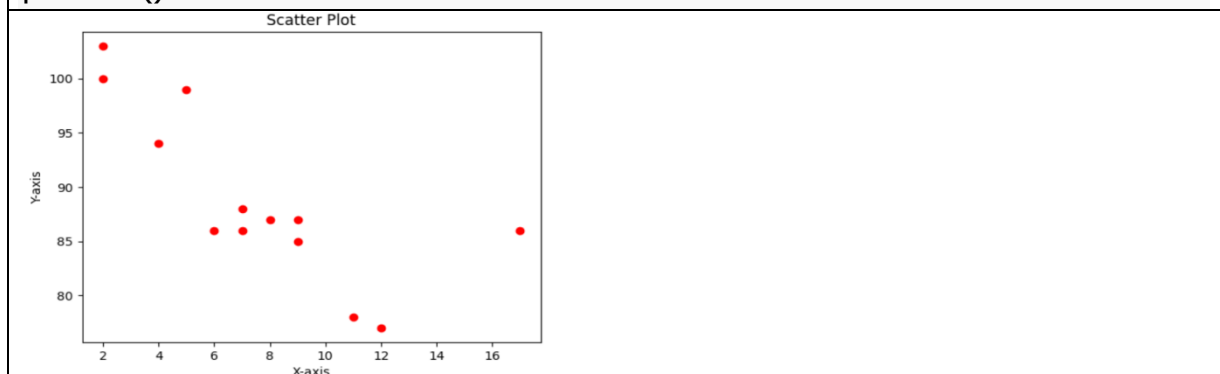
3) Histogram

```
import matplotlib.pyplot as plt
import numpy as np
data = np.random.randn(1000) # data=np.random.normal(0,1,1000)
# np.random.randn() is a NumPy function that returns samples from the "standard
# normal" distribution. Mean ( $\mu$ ) = 0 & Standard deviation ( $\sigma$ ) = 1
plt.hist(data, bins=30, color='blue', edgecolor='black')
# bins=30: No. of bins (or bars) in the histogram.
# color='blue': Color of the bars
# edgecolor='black': Color of the edges (borders) of the bars to black
plt.title('Histogram')
plt.xlabel('Value')
plt.ylabel('Frequency')
plt.show()
```



4) Scatter Plot

```
import matplotlib.pyplot as plt
x = [5, 7, 8, 7, 2, 17, 2, 9, 4, 11, 12, 9, 6]
y = [99, 86, 87, 88, 100, 86, 103, 87, 94, 78, 77, 85, 86]
plt.scatter(x, y, color='red')
plt.title('Scatter Plot')
plt.xlabel('X-axis')
plt.ylabel('Y-axis')
plt.show()
```



5) Pie Chart

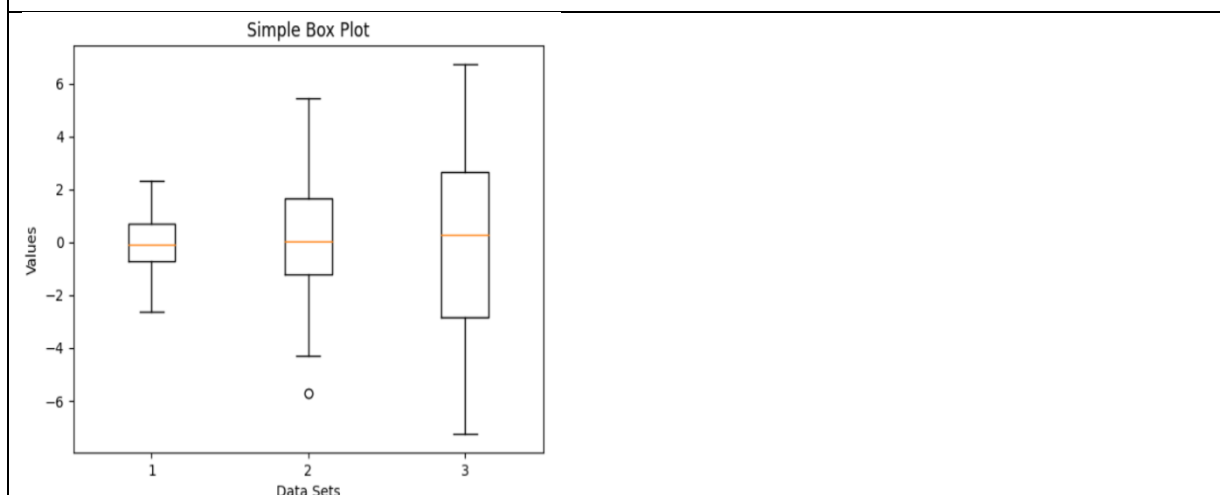
```
import matplotlib.pyplot as plt
sizes = [15, 30, 45, 10]
labels = ['A', 'B', 'C', 'D']
colors = ['gold', 'yellowgreen', 'lightcoral', 'lightskyblue']
plt.pie(sizes, labels=labels, colors=colors, autopct='%1.1f%%', shadow=True,
startangle=140)
plt.title('Pie Chart')
plt.axis('equal') # Equal aspect ratio ensures that pie is drawn as a circle.
plt.show()
```



6) Box Plot

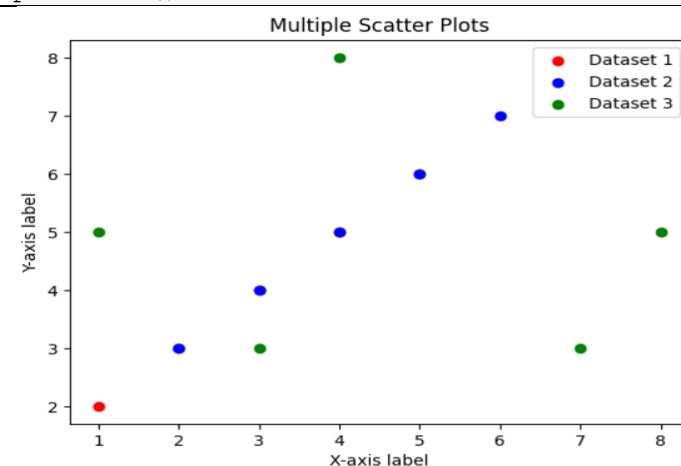
```
import matplotlib.pyplot as plt
import numpy as np

# Sample data
data = [np.random.normal(0, std, 100) for std in range(1, 4)]
# np.random.normal(0, std, 100) - np.random.normal(mean, std deviation, size)
# for std in range(1, 4) - list comprehension that iterates over a range of values ie, 1,2,3
plt.boxplot(data) # Create a box plot
# Add labels and title
plt.xlabel('Data Sets')
plt.ylabel('Values')
plt.title('Simple Box Plot')
# Show the plot
plt.show()
```



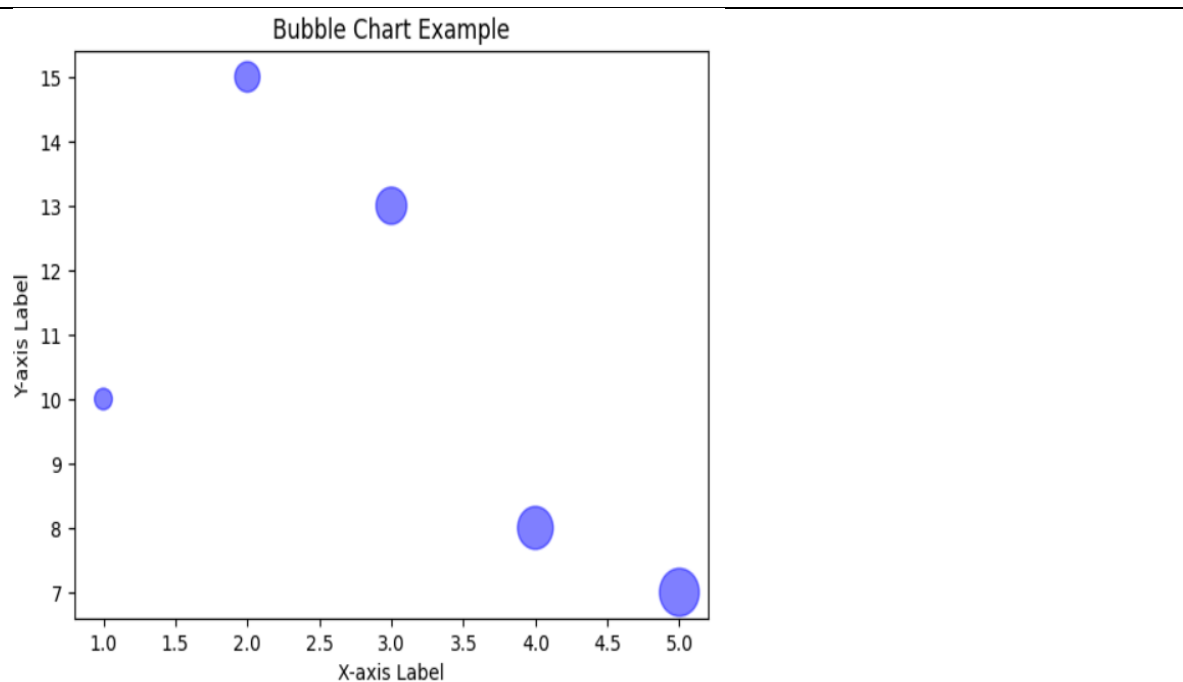
7) Scatter Multiple

```
import matplotlib.pyplot as plt
# Data for the first scatter plot
x1 = [1, 2, 3, 4, 5]
y1 = [2, 3, 4, 5, 6]
# Data for the second scatter plot
x2 = [2, 3, 4, 5, 6]
y2 = [3, 4, 5, 6, 7]
# Data for the third scatter plot
x3 = [1, 3, 4, 7, 8]
y3 = [5, 3, 8, 3, 5]
# Plotting the first scatter plot
plt.scatter(x1, y1, color='red', label='Dataset 1')
# Plotting the second scatter plot
plt.scatter(x2, y2, color='blue', label='Dataset 2')
# Plotting the third scatter plot
plt.scatter(x3, y3, color='green', label='Dataset 3')
# Adding labels and title
plt.xlabel('X-axis label')
plt.ylabel('Y-axis label')
plt.title('Multiple Scatter Plots')
plt.legend() # Adding a legend
plt.show()
```



8) Bubble Chart - A type of scatter plot where each point has an additional third dimension represented by the size of the bubble

```
# Creating a bubble chart
import matplotlib.pyplot as plt
# Sample data
x = [1, 2, 3, 4, 5]
y = [10, 15, 13, 8, 7]
sizes = [100, 200, 300, 400, 500] # Bubble sizes
plt.scatter(x, y, s=sizes, alpha=0.5, c='blue')
# Adding labels and title
plt.xlabel('X-axis Label')
plt.ylabel('Y-axis Label')
plt.title('Bubble Chart Example')
plt.show()
```

#Bubble chart

import matplotlib.pyplot as plt

Sample data with an additional dimension for colors

x = [1, 2, 3, 4, 5]

y = [10, 15, 13, 8, 7]

sizes = [100, 200, 300, 400, 500]

colors = [20, 30, 40, 50, 60] # Color dimension

Creating a bubble chart with varying colors

plt.scatter(x, y, s=sizes, c=colors, alpha=0.6, cmap='viridis')

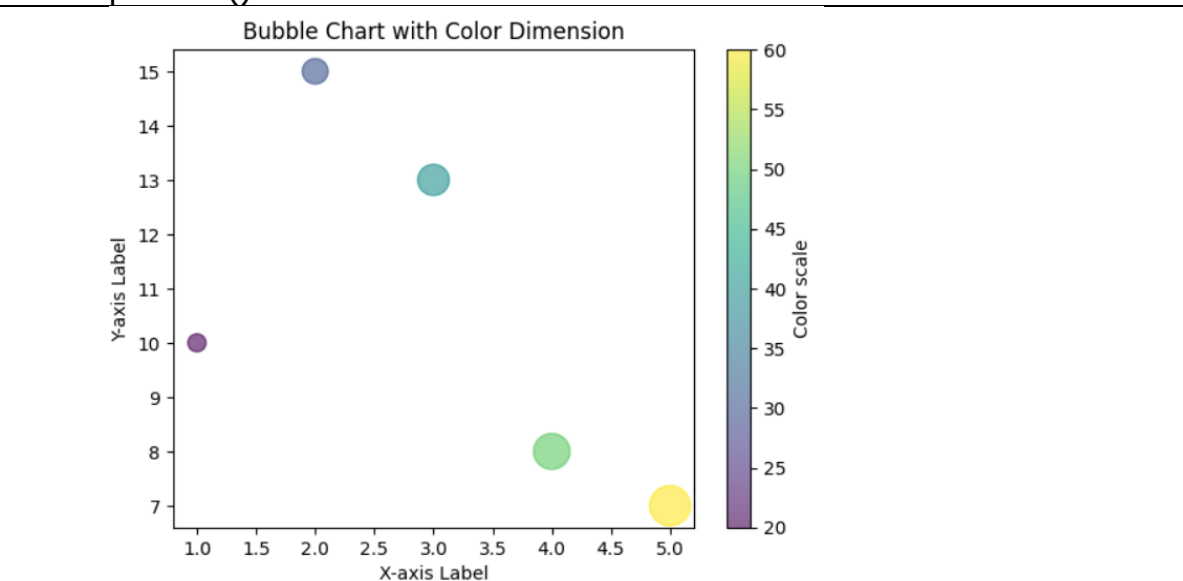
plt.colorbar(label='Color scale') # Adding a color bar

plt.xlabel('X-axis Label')

plt.ylabel('Y-axis Label')

plt.title('Bubble Chart with Color Dimension')

plt.show()

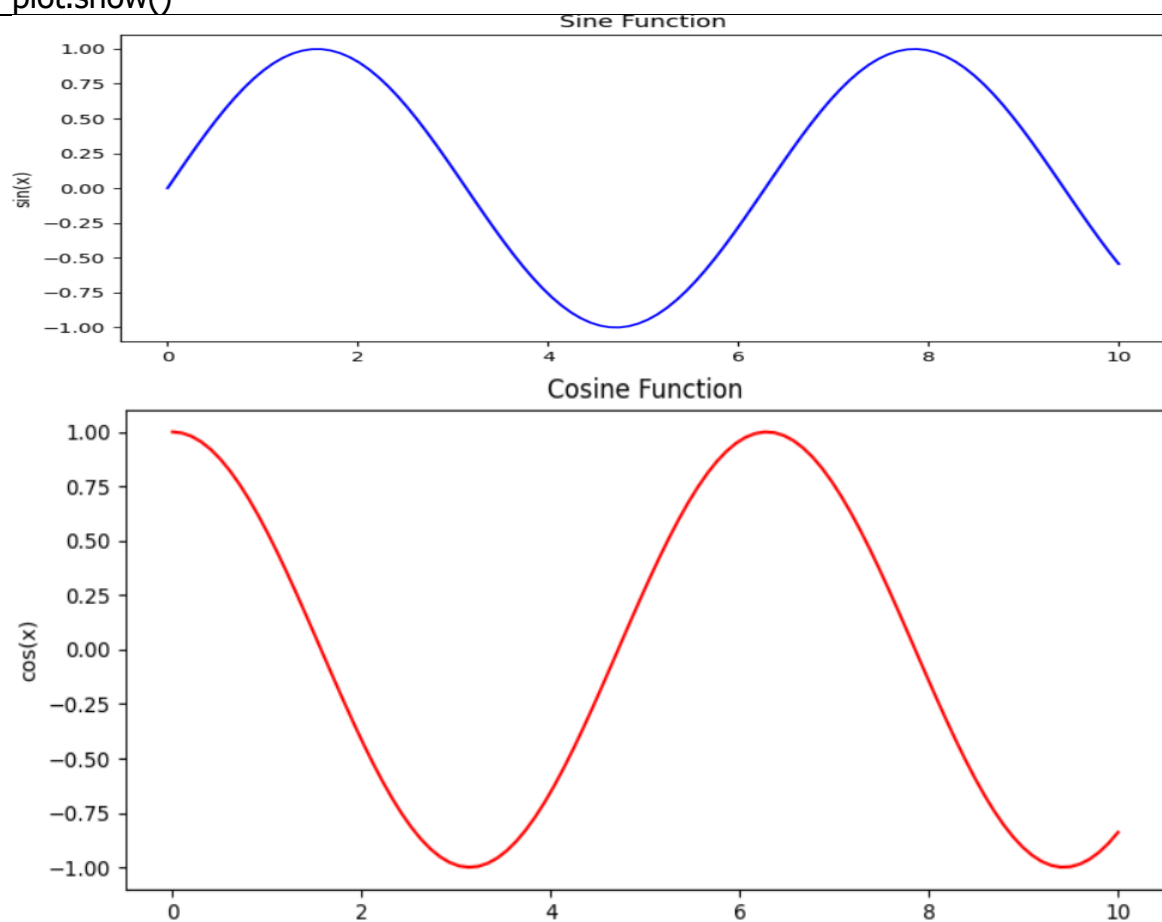


9) Subplots

```
import matplotlib.pyplot as plt
import numpy as np

# Sample data
x = np.linspace(0, 10, 100)
y1 = np.sin(x)
y2 = np.cos(x)
# Create a figure with 2 subplots (vertically stacked)
fig, axs = plt.subplots(2, 1, figsize=(8, 8))

# First subplot
axs[0].plot(x, y1, 'b')
axs[0].set_title('Sine Function')
axs[0].set_ylabel('sin(x)')
# Second subplot
axs[1].plot(x, y2, 'r')
axs[1].set_title('Cosine Function')
axs[1].set_ylabel('cos(x)')
axs[1].set_xlabel('x')
plt.tight_layout() # Adjust layout to prevent overlap
plt.show()
```



PROGRAM 7 : Sarah bought a new car in 2001 for \$24000. The dollar value of her car changed each year as shown in the table below.

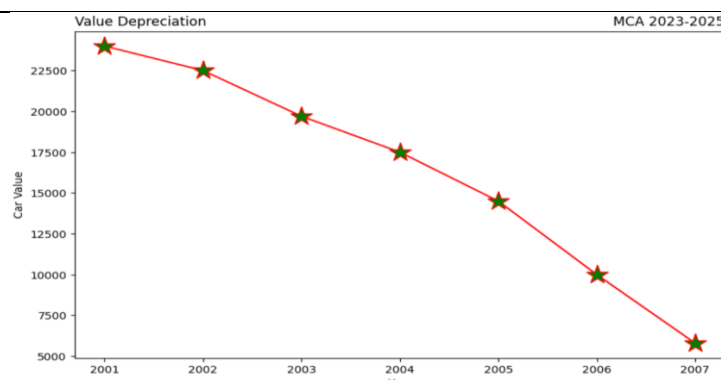
Value of Sarah's Car

Year	Value
2001	\$24,000
2002	\$22,500
2003	\$19,700
2004	\$17,500
2005	\$14,500
2006	\$10,000
2007	\$ 5,800

represent the following information using a line graph with following style properties

- X-axis – year
- Y-axis – car value
- Title – value depreciation (left aligned)
- Line style dash dot & line color should be red
- Point using * symbol with green color & size 20

```
import matplotlib.pyplot as plt
years = [2001, 2002, 2003, 2004, 2005, 2006, 2007]
car_values = [24000, 22500, 19700, 17500, 14500, 10000, 5800]
plt.figure(figsize=(10, 6))
plt.subplot(111)
plt.plot(years, car_values, linestyle='-', color='red', marker='*', markersize=20,
markerfacecolor='green')
plt.title(" \nMCA 2023-2025", loc="right")
plt.title("Value Depreciation", loc="left")
plt.xlabel("Year")
plt.ylabel("Car Value")
plt.show()
```



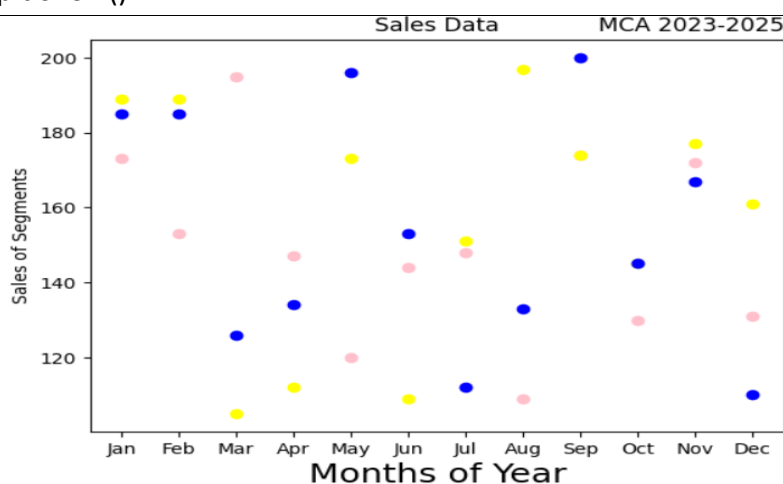
PROGRAM 8: Create scatter plot for the below data (use scatter function)

Product	Jan	Feb	Mar	Apr	May	Jun	Jul	Aug	Sep	Oct	Nov	Dec
Affordable Segment	173	153	195	147	120	144	148	109	174	130	172	131
Luxury Segment	189	189	105	112	173	109	151	197	174	145	177	161
Super Luxury Segment	185	185	126	134	196	153	112	133	200	145	167	110

Create scatter plot for each segment with following properties within one graph

- X-axis – months of year with font size 18
- Y-axis – sales of segments
- Title – sales data
- Color for affordable segment- pink
- Color for luxury segment – yellow
- Color for super luxury segment - blue

```
import matplotlib.pyplot as plt
import numpy as np
month = np.array(['Jan','Feb','Mar','Apr','May','Jun','Jul','Aug','Sep','Oct','Nov','Dec'])
AS = np.array([173,153,195,147,120,144,148,109,174,130,172,131])
LS = np.array([189,189,105,112,173,109,151,197,174,145,177,161])
SLS = np.array([185,185,126,134,196,153,112,133,200,145,167,110])
plt.xlabel('Months of Year', fontsize=18)
plt.ylabel('Sales of Segments')
plt.title('Sales Data')
plt.title('MCA 2023-2025', loc='right')
plt.scatter(month,AS, label='Affordable Segment', color='pink')
plt.scatter(month,LS, label='Luxury Segment', color='yellow')
plt.scatter(month,SLS, label='Super Luxury Segment', color='blue')
plt.show()
```



PROGRAM 9: Following table gives the daily sales of the following items in a shop.

Day	Mon	Tues	Wed	Thurs	Fri
Drinks	300	450	150	400	650
Food	400	500	350	300	500

Use subplot function to draw the line graphs with grids (color as blue & line style dotted) for the above information as 2 separate graphs in 2 rows

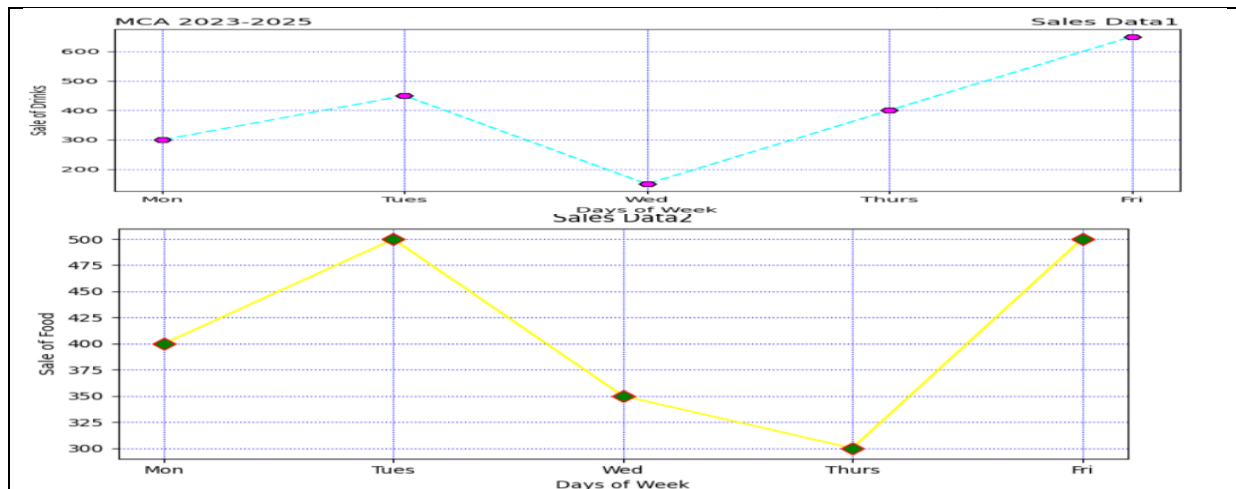
a) Properties for the graph1:

- X label – days of week
- Y label – Sale of drinks
- Title – sales data1 (right aligned)
- Line – dotted with cyan color
- Points – hexagon shape with color magenta & outline black

b) Properties for the graph2:

- X label – days of week
- Y label – Sale of food
- Title – sales data2 (center aligned)
- Line – dashed with yellow color
- Points – diamond shape with color green & outline red

```
import matplotlib.pyplot as plt
days = ['Mon', 'Tues', 'Wed', 'Thurs', 'Fri']
drinks_sales = [300, 450, 150, 400, 650]
food_sales = [400, 500, 350, 300, 500]
fig, axs = plt.subplots(2, 1, figsize=(8, 8))
axs[0].plot(days, drinks_sales, linestyle='--', color='cyan', marker='H',
markersize=8, markerfacecolor='magenta', markeredgecolor='black')
axs[0].set_xlabel('Days of Week')
axs[0].set_ylabel('Sale of Drinks')
axs[0].set_title('Sales Data1', loc='right')
axs[0].set_title('MCA 2023-2025', loc='left')
axs[0].grid(True, color='blue', linestyle='dotted')
axs[1].plot(days, food_sales, linestyle='-', color='yellow', marker='D', markersize=8,
markerfacecolor='green', markeredgecolor='red')
axs[1].set_xlabel('Days of Week')
axs[1].set_ylabel('Sale of Food')
axs[1].set_title('Sales Data2', loc='center')
axs[1].grid(True, color='blue', linestyle='dotted')
plt.tight_layout()
plt.show()
```



Pandas

Pandas is a Python library used for working with data sets. It has functions for analyzing, cleaning, exploring, and manipulating data.

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It has functions for analyzing, cleaning, exploring, and manipulating data. Pandas allow us to analyze big data and make conclusions based on statistical theories.

Pandas can clean messy data sets, and make them readable and relevant. Relevant data is very important in data science

A Pandas DataFrame is a 2 dimensional data structure, like a 2 dimensional array, or a table with rows and columns.

PROGRAM 10: Write programs to perform basic operations using pandas

```
#Import necessary modules
import numpy as np
import pandas as pd

#Creating a dataframe using List: DataFrame can be created using
#a single list or a list of lists.
data = {'Name': ['Tom', 'nick', 'krish', 'jack'], 'Age': [20, 21, 19, 18]}
df = pd.DataFrame(data) # Convert the dictionary into DataFrame
print(df)
```

	Name	Age
0	Tom	20
1	nick	21
2	krish	19
3	jack	18

```
#Dealing with Rows and Columns
# Define a dictionary containing employee data
data = {'Name': ['Jai', 'Princi', 'Gaurav', 'Anuj'],
        'Age': [27, 24, 22, 32],
        'Address': ['Delhi', 'Kanpur', 'Allahabad', 'Kannauj'],
        'Qualification': ['Msc', 'MA', 'MCA', 'Phd']}
df = pd.DataFrame(data) # Convert the dictionary into DataFrame
print(df[['Name', 'Qualification']]) # select two columns
```

	Name	Qualification
0	Jai	Msc
1	Princi	MA
2	Gaurav	MCA
3	AnujPhd	

```
data = pd.read_csv("/content/empl.csv")
```

```
data.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 33 entries, 0 to 32
Data columns (total 9 columns):
#   Column                Non-Null Count  Dtype
---  -
0   EMPLOYEE_ID           22 non-null    float64
1   NAME                  22 non-null    object
2   EMAIL                 23 non-null    object
3   PHONE_NUMBER          22 non-null    object
4   HIRE_DATE              22 non-null    object
5   DESIGNATION            22 non-null    object
6   SALARY                 22 non-null    float64
7   MANAGER_ID             22 non-null    object
8   DEPARTMENT_ID          22 non-null    float64
dtypes: float64(3), object(6)
memory usage: 2.4+ KB
```

```
data.head()
```

	EMPLOYEE_ID	NAME	EMAIL	PHONE_NUMBER	HIRE_DATE	DESIGNATION	SALARY	MANAGER_ID	DEPARTMENT_ID
0	198.0	Donald	DOCONNEL	650.507.9833	21-Jun-07	SH_CLERK	12600.0	124	50.0
1	199.0	Douglas	DGRANT	650.507.9844	13-Jan-08	SH_CLERK	12600.0	124	50.0
2	200.0	Jennifer	JWHALEN	515.123.4444	17-Sep-03	AD_ASST	14400.0	101	10.0
3	201.0	Michael	MHARTSTE	515.123.5555	17-Feb-04	MK_MAN	13000.0	100	20.0
4	202.0	Pat	PFAY	603.123.6666	17-Aug-05	MK_REP	16000.0	201	20.0

```
# retrieving columns by indexing operator
```

```
first = data.head()
```

```
print(first)
```

EMPLOYEE_ID	NAME	EMAIL	PHONE_NUMBER	HIRE_DATE	DESIGNATION	\
0	198.0	Donald	DOCONNEL	650.507.9833	21-Jun-07	SH_CLERK
1	199.0	Douglas	DGRANT	650.507.9844	13-Jan-08	SH_CLERK
2	200.0	Jennifer	JWHALEN	515.123.4444	17-Sep-03	AD_ASST
3	201.0	Michael	MHARTSTE	515.123.5555	17-Feb-04	MK_MAN
4	202.0	Pat	PFAY	603.123.6666	17-Aug-05	MK_REP

	SALARY	MANAGER_ID	DEPARTMENT_ID
0	12600.0	124	50.0
1	12600.0	124	50.0
2	14400.0	101	10.0
3	13000.0	100	20.0
4	16000.0	201	20.0

```
import pandas as pd
import numpy as np
arr=np.array([10,15,18,22])
s = pd.Series(arr)
print(s)
```

```
0    10
1    15
2    18
3    22
dtype: int64
```

```
arr=np.array(['a','b','c','d'])
s=pd.Series(arr, index=['first','second','third','fourth'])
print(s)
```

```
first      a
second     b
third      c
fourth     d
dtype: object
```

```
s=pd.Series(50, index=[0, 1, 2, 3, 4])
print (s)
```

```
0    50
1    50
2    50
3    50
4    50
dtype: int64
```

#Creating a series from a Dictionary

```
d={'Name': 'Deepthi', 'Class' : 'MCA', 'year' : 2014}
s=pd.Series(d)
print(s)
```

```
Name      Deepthi
Class      MCA
year      2014
dtype: object
```

#To Add & Rename a column in data frame

```
s = pd.Series([10,15,18,22])
df=pd.DataFrame(s)
df.columns=['List1']
#To Rename the default column of Data Frame as List1
df['List2']=20
#To create a new column List2 with all values as 20
df['List3']=df['List1']+df['List2']
#Add Column1 and Column2 and store in New column List3
print(df)
```

```
List1  List2  List3
0      10     20    30
1      15     20    35
2      18     20    38
3      22     20    42
```



```

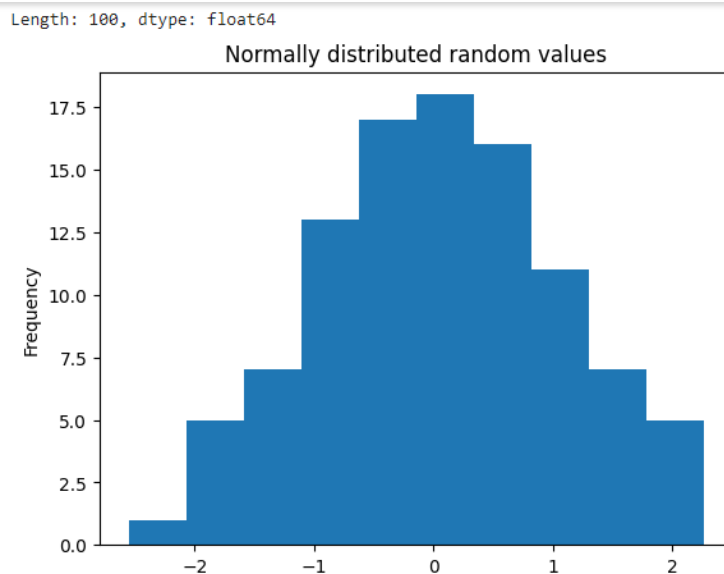
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
np.random.seed(0)
values = np.random.randn(100)
s = pd.Series(values) # generate a pandas series
print("series\n",s)
s.plot(kind='hist', title='Normally distributed random values')
plt.show()

```

```

series
0      1.764052
1      0.400157
2      0.978738
3      2.240893
4      1.867558
...
95      0.706573
96      0.010500
97      1.785870
98      0.126912
99      0.401989
Length: 100, dtype: float64

```



Describe -Descriptive statistics (mean, standard deviation, number of observations, minimum, maximum, and quartiles) of numerical columns can be calculated using the describe() method, which returns a pandas dataframe of descriptive statistics.

s.describe()

```

count      100.000000
mean         0.059808
std          1.012960
min         -2.552990
25%         -0.643857
50%          0.094096
75%          0.737077
max          2.269755
dtype: float64

```

```
df = pd.DataFrame({'A': [1, 2, 1, 4, 3],
'B': [12, 14, 11, 16, 18],
'C': ['a', 'a', 'b', 'a', 'b']})
df
```

	A	B	C
0	1	12	a
1	2	14	a
2	1	11	b
3	4	16	a
4	3	18	b

```
df.describe()
```

	A	B
count	5.00000	5.000000
mean	2.20000	14.200000
std	1.30384	2.863564
min	1.00000	11.000000
25%	1.00000	12.000000
50%	2.00000	14.000000
75%	3.00000	16.000000
max	4.00000	18.000000

```
#Display All records
import pandas
iris = pandas.read_csv('/content/Iris_new.csv')
print(iris)
```

output

	Id	SepalLength	SepalWidth	PetalLength	PetalWidth	Species
0	1	5.1	3.5	1.4	0.2	Iris-setosa
1	2	4.9	3.0	1.4	0.2	Iris-setosa
2	3	4.7	3.2	1.3	0.2	Iris-setosa
3	4	4.6	3.1	1.5	0.2	Iris-setosa
4	5	5.0	3.6	1.4	0.2	Iris-setosa
5	6	5.4	3.9	1.7	0.4	Iris-setosa
6	7	4.6	3.4	1.4	0.3	Iris-setosa
7	8	5.0	3.4	1.5	0.2	Iris-setosa
8	9	4.4	2.9	1.4	0.2	Iris-setosa

9	10	4.9	3.1	1.5	0.1	Iris-
setosa						
10	11	7.0	3.2	4.7	1.4	Iris-
versicolor						
11	12	6.4	3.2	4.5	1.5	Iris-
versicolor						
12	13	6.9	3.1	4.9	1.5	Iris-
versicolor						
13	14	5.5	2.3	4.0	1.3	Iris-
versicolor						
14	15	6.5	2.8	4.6	1.5	Iris-
versicolor						
15	16	5.7	2.8	4.5	1.3	Iris-
versicolor						
16	17	6.3	3.3	4.7	1.6	Iris-
versicolor						
17	18	4.9	2.4	3.3	1.0	Iris-
versicolor						
18	19	6.6	2.9	4.6	1.3	Iris-
versicolor						
19	20	5.2	2.7	3.9	1.4	Iris-
versicolor						
20	21	6.3	3.3	6.0	2.5	Iris-
virginica						
21	22	5.8	2.7	5.1	1.9	Iris-
virginica						
22	23	7.1	3.0	5.9	2.1	Iris-
virginica						
23	24	6.3	2.9	5.6	1.8	Iris-
virginica						
24	25	6.5	3.0	5.8	2.2	Iris-
virginica						
25	26	7.6	3.0	6.6	2.1	Iris-
virginica						
26	27	4.9	2.5	4.5	1.7	Iris-
virginica						
27	28	7.3	2.9	6.3	1.8	Iris-
virginica						
28	29	6.7	2.5	5.8	1.8	Iris-
virginica						
29	30	7.2	3.6	6.1	2.5	Iris-
virginica						

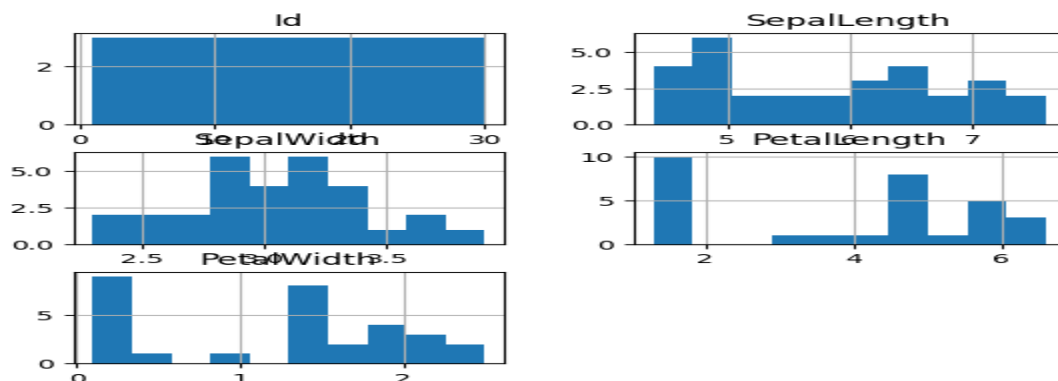
```
#display top 5 records
import pandas as pd
iris = pd.read_csv('/content/Iris_new.csv')
print(iris.head())
```

```
Output
Id  SepalLength  SepalWidth  PetalLength  PetalWidth  Species
0   1           5.1         3.5         1.4         0.2  Iris-setosa
1   2           4.9         3.0         1.4         0.2  Iris-setosa
2   3           4.7         3.2         1.3         0.2  Iris-setosa
3   4           4.6         3.1         1.5         0.2  Iris-setosa
4   5           5.0         3.6         1.4         0.2  Iris-setosa
```

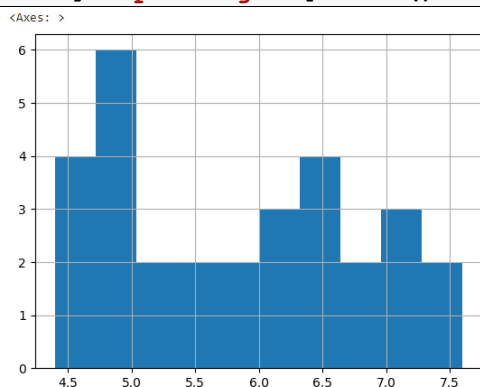
```
import pandas as pd
df = pd.DataFrame(pd.read_csv("/content/Iris_new.csv"))
# create histogram for numeric data
df.hist()
```

Output

```
array([[<Axes: title={'center': 'Id'}>, <Axes: title={'center': 'SepalLength'}>],
       [<Axes: title={'center': 'SepalWidth'}>, <Axes: title={'center': 'PetalLength'}>],
       [<Axes: title={'center': 'PetalWidth'}>, <Axes: >]], dtype=object)
```



```
iris['SepalLength'].hist()
```



Exploratory Data Analysis using pandas

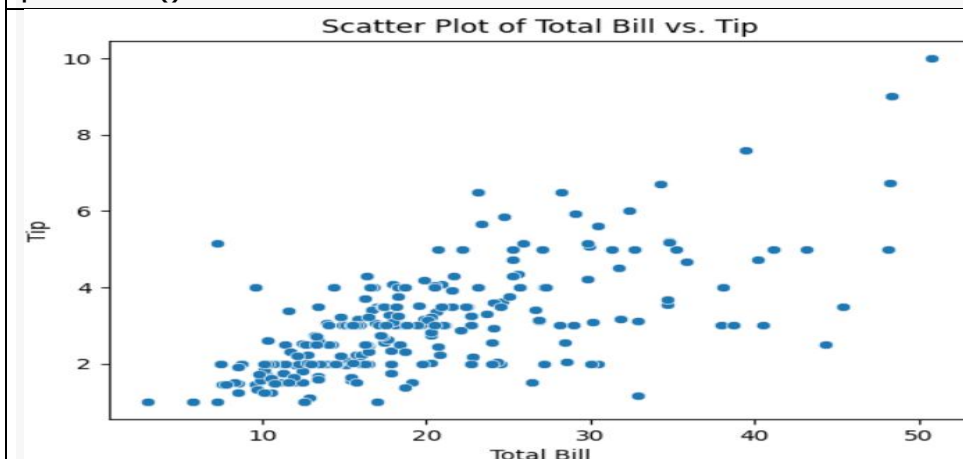
```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
# Load Housing Price dataset
data = pd.read_csv('/content/sample_data/california_housing_test.csv')
# Exploratory Data Analysis
data.info()
data.describe()
data.head()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 3000 entries, 0 to 2999
Data columns (total 9 columns):
#   Column                Non-Null Count  Dtype  
---  -
0   longitude              3000 non-null   float64
1   latitude               3000 non-null   float64
2   housing_median_age     3000 non-null   float64
3   total_rooms            3000 non-null   float64
4   total_bedrooms         3000 non-null   float64
5   population             3000 non-null   float64
6   households             3000 non-null   float64
7   median_income          3000 non-null   float64
8   median_house_value     3000 non-null   float64
dtypes: float64(9)
memory usage: 211.1 KB
```

	longitude	latitude	housing_median_age	total_rooms	total_bedrooms	population	households	median_income	median_house_value
0	-122.05	37.37	27.0	3885.0	661.0	1537.0	606.0	6.6085	342611.0
1	-118.30	34.26	43.0	1510.0	310.0	809.0	277.0	3.5990	153450.0
2	-117.81	33.78	27.0	3589.0	507.0	1484.0	495.0	5.7934	535350.0
3	-118.36	33.82	28.0	67.0	15.0	49.0	11.0	6.1359	341312.0
4	-119.67	36.33	19.0	1241.0	244.0	850.0	237.0	2.9375	181525.0

- **matplotlib** is a **foundational plotting library** for Python that provides extensive control over the appearance of plots. It's a general-purpose plotting library and is widely used for creating static, animated, and interactive visualizations.
- **seaborn** is a **statistical data visualization library** built on top of **matplotlib**. It provides a high-level interface for creating attractive and informative statistical graphics with simpler syntax.

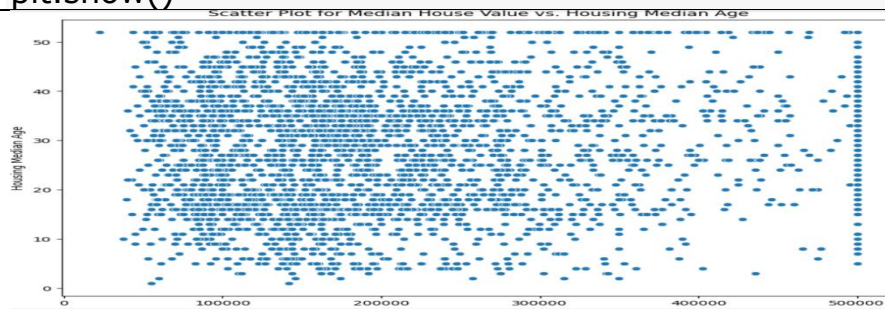
```
import seaborn as sns
import matplotlib.pyplot as plt
# Example Data
data = sns.load_dataset('tips')
# Scatter Plot
sns.scatterplot(x='total_bill', y='tip', data=data)
plt.title('Scatter Plot of Total Bill vs. Tip')
plt.xlabel('Total Bill')
plt.ylabel('Tip')
plt.show()
```



```

import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
# Load Housing Price dataset
data = pd.read_csv ('/content/sample_data/
california_housing_test.csv')
# Scatter Plot
plt.figure(figsize=(10, 8))
sns.scatterplot(x='median_house_value', y='housing_median_age',
data=data)
plt.title('Scatter Plot for Median House Value vs. Housing Median Age')
plt.xlabel('Median House Value')
plt.ylabel('Housing Median Age')
plt.show()

```



```

# Box Plot
plt.figure(figsize=(15, 10))
sns.boxplot(x='housing_median_age', y='median_house_value', data=data)
plt.title('Box Plot for Feature1 by Target')
plt.show()

```

Program 11: Dataset: <Breast_Cancer.csv>

- Conduct **exploratory data analysis** on the given dataset and report the details.
- Visualize** the analysis results using
 - scatter plot
 - histogram
 - box plot
- Implement the **k-NN classification algorithm** using the dataset. Try with **different k values and show the accuracy**.

#a) exploratory data analysis

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.datasets import load_breast_cancer
from sklearn.model_selection import train_test_split
from sklearn.neighbors import KNeighborsClassifier
from sklearn import metrics
```

Load the Breast Cancer dataset

```
data = load_breast_cancer()
df = pd.DataFrame(data.data, columns=data.feature_names)
df['target'] = data.target
print(df.head(5))
df.info()
```

```
mean radius mean texture mean perimeter mean area mean smoothness \
0    17.99    10.38    122.80    1001.0    0.11840
1    20.57    17.77    132.90    1326.0    0.08474
2    19.69    21.25    130.00    1203.0    0.10960
3    11.42    20.38    77.58    386.1    0.14250
4    20.29    14.34    135.10    1297.0    0.10030
```

```
mean compactness mean concavity mean concave points mean symmetry \
0    0.27760    0.3001    0.14710    0.2419
1    0.07864    0.0869    0.07017    0.1812
2    0.15990    0.1974    0.12790    0.2069
3    0.28390    0.2414    0.10520    0.2597
4    0.13280    0.1980    0.10430    0.1809
```

```
mean fractal dimension ... worst texture worst perimeter worst area \
0    0.07871 ...    17.33    184.60    2019.0
1    0.05667 ...    23.41    158.80    1956.0
2    0.05999 ...    25.53    152.50    1709.0
3    0.09744 ...    26.50    98.87    567.7
4    0.05883 ...    16.67    152.20    1575.0
```

```
worst smoothness worst compactness worst concavity worst concave points \
0    0.1622    0.6656    0.7119    0.2654
1    0.1238    0.1866    0.2416    0.1860
2    0.1444    0.4245    0.4504    0.2430
3    0.2098    0.8663    0.6869    0.2575
4    0.1374    0.2050    0.4000    0.1625
```

```
worst symmetry worst fractal dimension target
0    0.4601    0.11890    0
1    0.2750    0.08902    0
2    0.3613    0.08758    0
3    0.6638    0.17300    0
4    0.2364    0.07678    0
```

```
[5 rows x 31 columns]
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 569 entries, 0 to 568
Data columns (total 31 columns):
#   Column                                Non-Null Count  Dtype
---  -
0   mean radius                           569 non-null    float64
1   mean texture                           569 non-null    float64
2   mean perimeter                         569 non-null    float64
3   mean area                             569 non-null    float64
4   mean smoothness                       569 non-null    float64
5   mean compactness                      569 non-null    float64
6   mean concavity                        569 non-null    float64
7   mean concave points                   569 non-null    float64
8   mean symmetry                         569 non-null    float64
9   mean fractal dimension                569 non-null    float64
10  radius error                          569 non-null    float64
11  texture error                         569 non-null    float64
12  perimeter error                       569 non-null    float64
13  area error                           569 non-null    float64
14  smoothness error                     569 non-null    float64
15  compactness error                    569 non-null    float64
16  concavity error                      569 non-null    float64
17  concave points error                 569 non-null    float64
18  symmetry error                       569 non-null    float64
19  fractal dimension error              569 non-null    float64
20  worst radius                         569 non-null    float64
21  worst texture                        569 non-null    float64
22  worst perimeter                      569 non-null    float64
23  worst area                           569 non-null    float64
24  worst smoothness                     569 non-null    float64
25  worst compactness                    569 non-null    float64
26  worst concavity                      569 non-null    float64
27  worst concave points                 569 non-null    float64
28  worst symmetry                       569 non-null    float64
29  worst fractal dimension              569 non-null    float64
30  target                              569 non-null    int64
dtypes: float64(30), int64(1)
memory usage: 137.9 KB
```

```
print(df.describe)
<bound method NDFrame.describe of      mean radius  mean texture  mean perimeter  mean area
mean smoothness \
0      17.99      10.38      122.80      1001.0      0.11840
1      20.57      17.77      132.90      1326.0      0.08474
2      19.69      21.25      130.00      1203.0      0.10960
3      11.42      20.38      77.58      386.1      0.14250
4      20.29      14.34      135.10      1297.0      0.10030
..      ...      ...      ...      ...      ...
564     21.56     22.39     142.00     1479.0      0.11100
565     20.13     28.25     131.20     1261.0      0.09780
566     16.60     28.08     108.30      858.1      0.08455
567     20.60     29.33     140.10     1265.0      0.11780
568      7.76     24.54      47.92      181.0      0.05263

      mean compactness  mean concavity  mean concave points  mean symmetry \
0      0.27760      0.30010      0.14710      0.2419
1      0.07864      0.08690      0.07017      0.1812
2      0.15990      0.19740      0.12790      0.2069
3      0.28390      0.24140      0.10520      0.2597
4      0.13280      0.19800      0.10430      0.1809
..      ...      ...      ...      ...
564     0.11590      0.24390      0.13890      0.1726
565     0.10340      0.14400      0.09791      0.1752
566     0.10230      0.09251      0.05302      0.1590
```


567	0.27700	0.35140	0.15200	0.2397
568	0.04362	0.00000	0.00000	0.1587
mean fractal dimension ... worst texture worst perimeter worst area \				
0	0.07871	...	17.33	184.60 2019.0
1	0.05667	...	23.41	158.80 1956.0
2	0.05999	...	25.53	152.50 1709.0
3	0.09744	...	26.50	98.87 567.7
4	0.05883	...	16.67	152.20 1575.0
..
564	0.05623	...	26.40	166.10 2027.0
565	0.05533	...	38.25	155.00 1731.0
566	0.05648	...	34.12	126.70 1124.0
567	0.07016	...	39.42	184.60 1821.0
568	0.05884	...	30.37	59.16 268.6
worst smoothness worst compactness worst concavity \				
0	0.16220	0.66560	0.7119	
1	0.12380	0.18660	0.2416	
2	0.14440	0.42450	0.4504	
3	0.20980	0.86630	0.6869	
4	0.13740	0.20500	0.4000	
..	
564	0.14100	0.21130	0.4107	
565	0.11660	0.19220	0.3215	
566	0.11390	0.30940	0.3403	
567	0.16500	0.86810	0.9387	
568	0.08996	0.06444	0.0000	
worst concave points worst symmetry worst fractal dimension target				
0	0.2654	0.4601	0.11890	0
1	0.1860	0.2750	0.08902	0
2	0.2430	0.3613	0.08758	0
3	0.2575	0.6638	0.17300	0
4	0.1625	0.2364	0.07678	0
..
print(df.isnull().sum()) # Check for missing values print(df.describe()) # Summary statistics of the dataset # Target class distribution (Benign = 1, Malignant = 0) # Benign (1): Non-cancerous tumor Malignant (0): Cancerous tumor. print(df['target'].value_counts())				
mean radius	0			
mean texture	0			
mean perimeter	0			
mean area	0			
mean smoothness	0			
mean compactness	0			
mean concavity	0			
mean concave points	0			
mean symmetry	0			
mean fractal dimension	0			
radius error	0			
texture error	0			
perimeter error	0			
area error	0			
smoothness error	0			
compactness error	0			
concavity error	0			
concave points error	0			
symmetry error	0			
fractal dimension error	0			
worst radius	0			
worst texture	0			
worst perimeter	0			

```

worst area          0
worst smoothness    0
worst compactness   0
worst concavity      0
worst concave points 0
worst symmetry       0
worst fractal dimension 0
target              0
dtype: int64

      mean radius mean texture mean perimeter mean area \
count  569.000000  569.000000   569.000000  569.000000
mean   14.127292  19.289649   91.969033  654.889104
std     3.524049   4.301036   24.298981  351.914129
min     6.981000   9.710000   43.790000  143.500000
25%    11.700000  16.170000   75.170000  420.300000
50%    13.370000  18.840000   86.240000  551.100000
75%    15.780000  21.800000  104.100000  782.700000
max    28.110000  39.280000  188.500000 2501.000000

      mean smoothness mean compactness mean concavity mean concave points \
count  569.000000   569.000000   569.000000   569.000000
mean    0.096360    0.104341    0.088799    0.048919
std     0.014064    0.052813    0.079720    0.038803
min     0.052630    0.019380    0.000000    0.000000
25%     0.086370    0.064920    0.029560    0.020310
50%     0.095870    0.092630    0.061540    0.033500
75%     0.105300    0.130400    0.130700    0.074000
max     0.163400    0.345400    0.426800    0.201200

mean symmetry mean fractal dimension ... worst texture \
count  569.000000   569.000000 ... 569.000000
mean    0.181162    0.062798 ... 25.677223
std     0.027414    0.007060 ... 6.146258
min     0.106000    0.049960 ... 12.020000
25%     0.161900    0.057700 ... 21.080000
50%     0.179200    0.061540 ... 25.410000
75%     0.195700    0.066120 ... 29.720000
max     0.304000    0.097440 ... 49.540000

      worst perimeter worst area worst smoothness worst compactness \
count  569.000000  569.000000   569.000000   569.000000
mean   107.261213  880.583128    0.132369    0.254265
std    33.602542  569.356993    0.022832    0.157336
min    50.410000  185.200000    0.071170    0.027290
25%    84.110000  515.300000    0.116600    0.147200
50%    97.660000  686.500000    0.131300    0.211900
75%   125.400000 1084.000000    0.146000    0.339100
max   251.200000 4254.000000    0.222600    1.058000

      worst concavity worst concave points worst symmetry \
count  569.000000   569.000000   569.000000
mean    0.272188    0.114606    0.290076
std     0.208624    0.065732    0.061867
min     0.000000    0.000000    0.156500
25%     0.114500    0.064930    0.250400
50%     0.226700    0.099930    0.282200
75%     0.382900    0.161400    0.317900
max     1.252000    0.291000    0.663800

worst fractal dimension target
count  569.000000  569.000000
mean    0.083946  0.627417
std     0.018061  0.483918
min     0.055040  0.000000
25%     0.071460  0.000000
50%     0.080040  1.000000
75%     0.092080  1.000000

```

```
max          0.207500  1.000000
```

```
[8 rows x 31 columns]
```

```
target
```

```
1    357
```

```
0    212
```

```
Name: count, dtype: int64
```

#b) **Visualization**

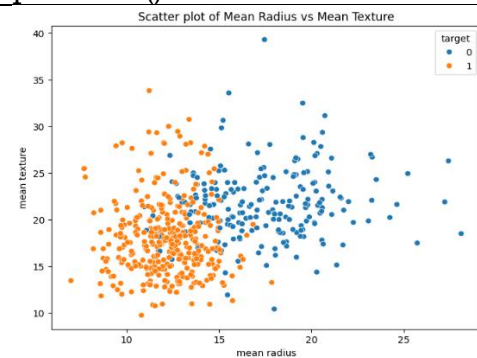
```
# (i) Scatter plot
```

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

```
sns.scatterplot(x='mean radius', y='mean texture', hue='target', data=df)
```

```
plt.title("Scatter plot of Mean Radius vs Mean Texture")
```

```
plt.show()
```



```
# (ii) Histogram
```

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

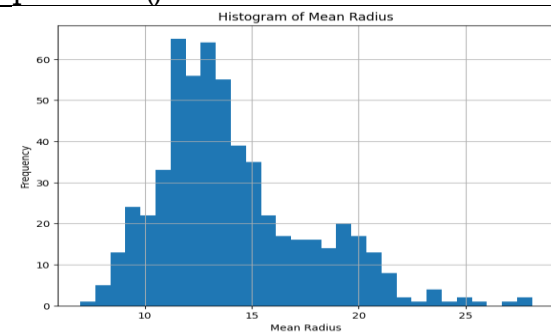
```
df['mean radius'].hist(bins=30)
```

```
plt.title("Histogram of Mean Radius")
```

```
plt.xlabel("Mean Radius")
```

```
plt.ylabel("Frequency")
```

```
plt.show()
```



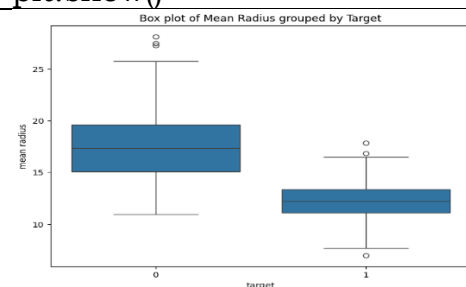
```
# (iii) Box plot
```

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

```
sns.boxplot(x='target', y='mean radius', data=df)
```

```
plt.title("Box plot of Mean Radius grouped by Target")
```

```
plt.show()
```



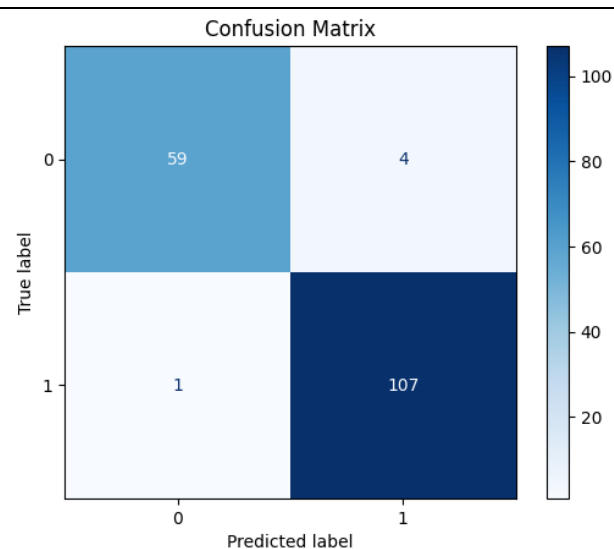
#c) Implementing k-NN Classification

```
# Splitting the dataset into training and testing sets
X = df.drop('target', axis=1) # Features
y = df['target'] # Target
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3,
random_state=42)
# Testing k-NN with different values of k
k_values = [3, 5, 7, 9]
for k in k_values:
    knn = KNeighborsClassifier(n_neighbors=k)
    knn.fit(X_train, y_train)
    y_pred = knn.predict(X_test)
accuracy = metrics.accuracy_score(y_test, y_pred) # Calculate accuracy
print(f'Accuracy with k={k}: {accuracy:.4f}')
```

```
Accuracy with k=3: 0.9415
Accuracy with k=5: 0.9591
Accuracy with k=7: 0.9649
Accuracy with k=9: 0.9708
```

#confusion matrix

```
from sklearn.metrics import confusion_matrix, ConfusionMatrixDisplay
# Confusion matrix
cm = confusion_matrix(y_test, y_pred, labels=knn.classes_)
cm_display = ConfusionMatrixDisplay(confusion_matrix=cm,
display_labels=knn.classes_)
cm_display.plot(cmap="Blues")
plt.title("Confusion Matrix")
plt.show()
```



Program 12: Dataset: <Wine_Quality.csv>

- Conduct **exploratory data analysis** on the given dataset and report the details.
- Visualize** the analysis results using (i) scatter plot (ii) histogram & (iii) box plot.
- Implement the **k-NN classification** algorithm using the dataset. Try with different K values and show the accuracy.

a) exploratory data analysis

import pandas as pd

Load the Wine Quality dataset

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

print(df.head()) **# Check the first few rows**

print(df.isnull().sum()) **# Check for missing values**

print(df.describe()) **# Summary statistics of the dataset**

Target variable distribution (if the dataset contains quality classes)

print(df['quality'].value_counts())

```

type  fixed acidity  volatile acidity  citric acid  residual sugar  \
0  white           7.0             0.27         0.36          20.7
1  white           6.3             0.30         0.34           1.6
2  white           8.1             0.28         0.40           6.9
3  white           7.2             0.23         0.32           8.5
4  white           7.2             0.23         0.32           8.5

chlorides  free sulfur dioxide  total sulfur dioxide  density  pH  \
0    0.045             45.0             170.0  1.0010  3.00
1    0.049             14.0             132.0  0.9940  3.30
2    0.050             30.0             97.0   0.9951  3.26
3    0.058             47.0             186.0  0.9956  3.19

```

```

4    0.058      47.0      186.0  0.9956  3.19

sulphates alcohol quality
0    0.45      8.8      6
1    0.49      9.5      6
2    0.44     10.1      6
3    0.40      9.9      6
4    0.40      9.9      6
type      0
fixed acidity      10
volatile acidity   8
citric acid        3
residual sugar     2
chlorides          2
free sulfur dioxide 0
total sulfur dioxide 0
density           0
pH                9
sulphates         4
alcohol           0
quality           0
dtype: int64
fixed acidity volatile acidity citric acid residual sugar \
count  6487.000000    6489.000000  6494.000000   6495.000000
mean    7.216579      0.339691    0.318722    5.444326
std     1.296750      0.164649    0.145265    4.758125
min     3.800000      0.080000    0.000000    0.600000
25%     6.400000      0.230000    0.250000    1.800000
50%     7.000000      0.290000    0.310000    3.000000
75%     7.700000      0.400000    0.390000    8.100000
max     15.900000     1.580000    1.660000   65.800000

chlorides free sulfur dioxide total sulfur dioxide density \
count  6495.000000    6497.000000    6497.000000  6497.000000
mean    0.056042      30.525319    115.744574    0.994697
std     0.035036      17.749400    56.521855    0.002999
min     0.009000      1.000000     6.000000    0.987110
25%     0.038000      17.000000    77.000000    0.992340
50%     0.047000      29.000000   118.000000    0.994890
75%     0.065000      41.000000   156.000000    0.996990
max     0.611000     289.000000   440.000000    1.038980

pH sulphates alcohol quality
count  6488.000000  6493.000000  6497.000000  6497.000000
mean    3.218395    0.531215   10.491801    5.818378
std     0.160748    0.148814    1.192712    0.873255
min     2.720000    0.220000    8.000000    3.000000
25%     3.110000    0.430000    9.500000    5.000000
50%     3.210000    0.510000   10.300000    6.000000
75%     3.320000    0.600000   11.300000    6.000000
max     4.010000    2.000000   14.900000    9.000000
quality
6    2836
5    2138
7    1079
4    216
8    193
3     30
9      5
Name: count, dtype: int64

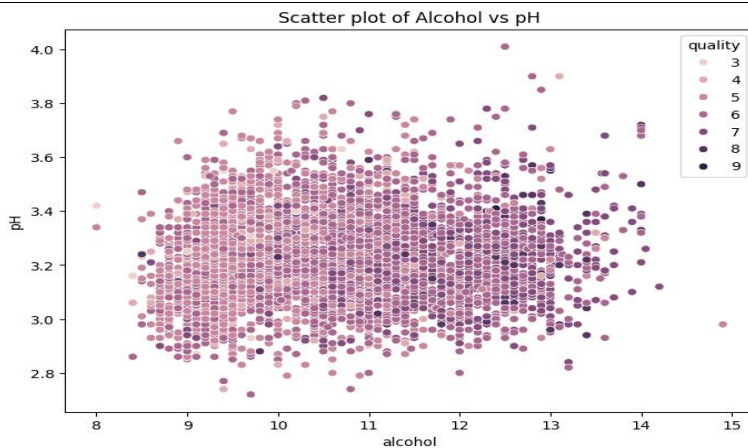
```

```
# b) Visualization
```

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

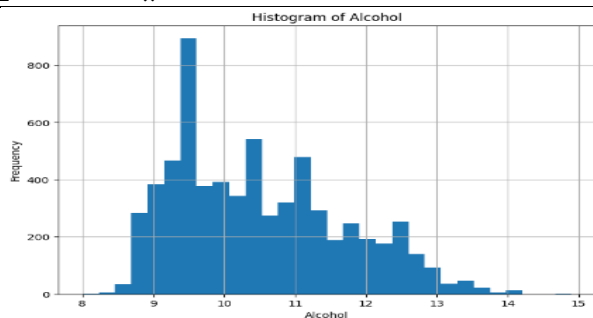
```
# (i) Scatter plot
```

```
plt.figure(figsize=(8, 6))
sns.scatterplot(x='alcohol', y='pH', hue='quality', data=df)
plt.title("Scatter plot of Alcohol vs pH")
plt.show()
```



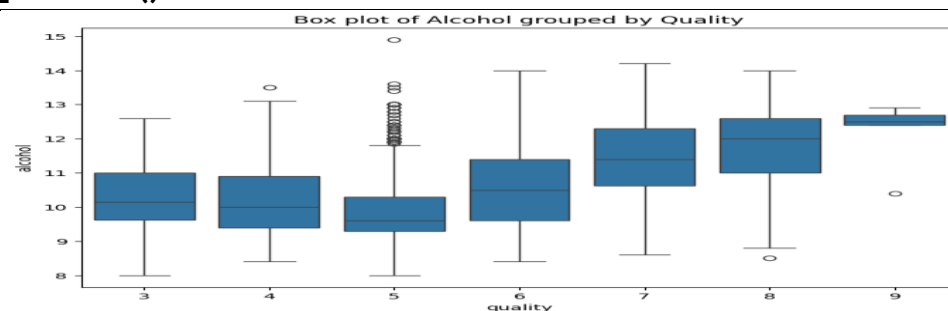
(ii) Histogram

```
plt.figure(figsize=(8, 6))
df['alcohol'].hist(bins=30)
plt.title("Histogram of Alcohol")
plt.xlabel("Alcohol")
plt.ylabel("Frequency")
plt.show()
```



Box plot

```
plt.figure(figsize=(8, 6))
sns.boxplot(x='quality', y='alcohol', data=df)
plt.title("Box plot of Alcohol grouped by Quality")
plt.show()
```



c) Implementing k-NN Classification

```
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import StandardScaler
from sklearn.neighbors import KNeighborsClassifier
from sklearn.metrics import accuracy_score

df_cleaned = df.dropna()# Drop rows or columns with missing values (if any)
```

```
# Define features (X) and target (y)
X = df_cleaned.drop(['quality', 'type'], axis=1) # Exclude 'quality' and 'type' for features
y = df_cleaned['quality'] # 'quality' is the target variable
# Split the data into training and testing sets (80% train, 20% test)
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2,
random_state=42)
scaler = StandardScaler()# Standardize the features for better performance of k-NN
X_train_scaled = scaler.fit_transform(X_train)
X_test_scaled = scaler.transform(X_test)
```

```
k_values = [1, 3, 5, 7, 9, 11, 13, 15]
for k in k_values:
    knn = KNeighborsClassifier(n_neighbors=k)
    knn.fit(X_train_scaled, y_train)
    # Predict on the test set
    y_pred = knn.predict(X_test_scaled)
    # Calculate accuracy
    accuracy = accuracy_score(y_test, y_pred)
    print(f'Accuracy with k={k}: {accuracy:.4f}')
```

```
Accuracy with k=1: 0.6226
Accuracy with k=3: 0.5538
Accuracy with k=5: 0.5584
Accuracy with k=7: 0.5313
Accuracy with k=9: 0.5445
Accuracy with k=11: 0.5491
Accuracy with k=13: 0.5483
Accuracy with k=15: 0.5429
```

```
k_values_input = input("Enter k values : ")# user to input k values from the console
# Convert the input string into a list of integers
k_values = [int(k.strip()) for k in k_values_input.split(',')]
for k in k_values:
    knn = KNeighborsClassifier(n_neighbors=k)
    knn.fit(X_train_scaled, y_train)
    y_pred = knn.predict(X_test_scaled) # Predict on the test set
    accuracy = accuracy_score(y_test, y_pred) # Calculate accuracy
    print(f'Accuracy with k={k}: {accuracy:.4f}')
```

```
Enter k values separated by commas (e.g., 3,5,7): 3
Accuracy with k=3: 0.5538
```


Program 13: Dataset: <Breast_Cancer.csv>

- Conduct exploratory data analysis on the given dataset and report the details.
- Visualize the analysis results using (i) scatter plot (ii) histogram & (iii) box plot.
- Implement the **Naïve Bayes classification** algorithm using the dataset. Display the classification report with the accuracy.

#a) Conduct exploratory data analysis

```
import pandas as pd
df = pd.read_csv('/content/breast-cancer.csv') # Load the Breast Cancer dataset
print(df.head()) # first few rows
print(df.isnull().sum()) # Check for missing values
print(df.describe()) # Summary statistics of the dataset
# Target variable distribution (if the dataset contains target classes, such as 'diagnosis')
if 'diagnosis' in df.columns:
    print(df['diagnosis'].value_counts())
```

```
symmetry_worst      0
fractal_dimension_worst  0
dtype: int64
      id  radius_mean  texture_mean  perimeter_mean  area_mean \
count  5.690000e+02  569.000000    569.000000    569.000000  569.000000
mean   3.037183e+07  14.127292    19.289649     91.969033  654.889104
std    1.250206e+08  3.524049     4.301036    24.298981  351.914129
min    8.670000e+03  6.981000     9.710000    43.790000  143.500000
25%    8.692180e+05  11.700000    16.170000    75.170000  420.300000
50%    9.060240e+05  13.370000    18.840000    86.240000  551.100000
75%    8.813129e+06  15.780000    21.800000   104.100000  782.700000
```

```
max 9.113205e+08 28.110000 39.280000 188.500000 2501.000000
```

```
smoothness_mean compactness_mean concavity_mean concave points_mean \
count 569.000000 569.000000 569.000000 569.000000
mean 0.096360 0.104341 0.088799 0.048919
std 0.014064 0.052813 0.079720 0.038803
min 0.052630 0.019380 0.000000 0.000000
25% 0.086370 0.064920 0.029560 0.020310
50% 0.095870 0.092630 0.061540 0.033500
75% 0.105300 0.130400 0.130700 0.074000
max 0.163400 0.345400 0.426800 0.201200
```

```
symmetry_mean ... radius_worst texture_worst perimeter_worst \
count 569.000000 ... 569.000000 569.000000 569.000000
mean 0.181162 ... 16.269190 25.677223 107.261213
std 0.027414 ... 4.833242 6.146258 33.602542
min 0.106000 ... 7.930000 12.020000 50.410000
25% 0.161900 ... 13.010000 21.080000 84.110000
50% 0.179200 ... 14.970000 25.410000 97.660000
75% 0.195700 ... 18.790000 29.720000 125.400000
max 0.304000 ... 36.040000 49.540000 251.200000
```

```
area_worst smoothness_worst compactness_worst concavity_worst \
count 569.000000 569.000000 569.000000 569.000000
mean 880.583128 0.132369 0.254265 0.272188
std 569.356993 0.022832 0.157336 0.208624
min 185.200000 0.071170 0.027290 0.000000
25% 515.300000 0.116600 0.147200 0.114500
50% 686.500000 0.131300 0.211900 0.226700
75% 1084.000000 0.146000 0.339100 0.382900
max 4254.000000 0.222600 1.058000 1.252000
```

```
concave points_worst symmetry_worst fractal_dimension_worst
count 569.000000 569.000000 569.000000
mean 0.114606 0.290076 0.083946
std 0.065732 0.061867 0.018061
min 0.000000 0.156500 0.055040
25% 0.064930 0.250400 0.071460
50% 0.099930 0.282200 0.080040
75% 0.161400 0.317900 0.092080
max 0.291000 0.663800 0.207500
```

```
[8 rows x 31 columns]
```

```
diagnosis
```

```
B 357
```

```
M 212
```

```
Name: count, dtype: int64
```

```
# b) visualization
```

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

```
import seaborn as sns
```

```
# Scatter plot (example: comparing 'radius_mean' and 'texture_mean')
```

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

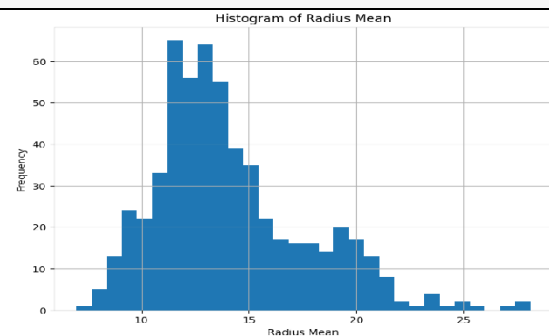
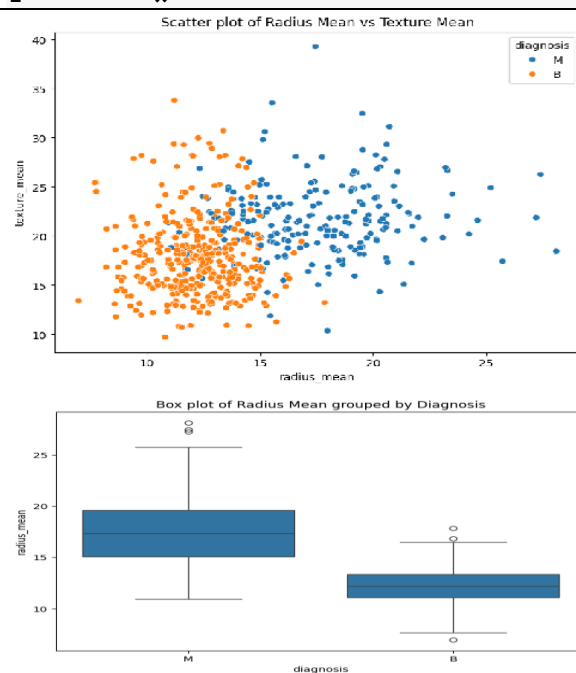
```
sns.scatterplot(x='radius_mean', y='texture_mean', hue='diagnosis', data=df)
```

```
plt.title("Scatter plot of Radius Mean vs Texture Mean")
```

```
plt.show()
```

```
# Histogram (example: distribution of 'radius_mean')
plt.figure(figsize=(8, 6))
df['radius_mean'].hist(bins=30)
plt.title("Histogram of Radius Mean")
plt.xlabel("Radius Mean")
plt.ylabel("Frequency")
plt.show()

# Box plot (example: distribution of 'radius_mean' grouped by 'diagnosis')
plt.figure(figsize=(8, 6))
sns.boxplot(x='diagnosis', y='radius_mean', data=df)
plt.title("Box plot of Radius Mean grouped by Diagnosis")
plt.show()
```



(c): Implementing Naïve Bayes Classification

```
from sklearn.model_selection import train_test_split
```

```
from sklearn.naive_bayes import GaussianNB
```

```
from sklearn.metrics import classification_report, accuracy_score
```

```
# df.dropna() - In Pandas to remove rows (or columns) that contain missing
# values from a DataFrame
```

```
df_cleaned = df.dropna()
```

```
# Define features (X) and target (y)
```

```
X = df_cleaned.drop(['diagnosis'], axis=1) # Exclude 'diagnosis' for features
```

```
y = df_cleaned['diagnosis'] # 'diagnosis' is the target variable
```

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

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

```
nb = GaussianNB() # Initialize the Naïve Bayes classifier
```

```
nb.fit(X_train, y_train) # Train the model
```

```
y_pred = nb.predict(X_test) # Predict on the test set
```

```
# Display the classification report and accuracy
print("Classification Report:\n", classification_report(y_test, y_pred))
accuracy = accuracy_score(y_test, y_pred)
print(f'Accuracy: {accuracy:.4f}')
```

Classification Report:

	precision	recall	f1-score	support
B	0.62	0.99	0.76	71
M	0.00	0.00	0.00	43
accuracy			0.61	114
macro avg	0.31	0.49	0.38	114
weighted avg	0.39	0.61	0.47	114

Accuracy: 0.6140

Program 14: Dataset: <Wine_Quality.csv>

- Conduct exploratory data analysis on the given dataset and report the details.
- Visualize the analysis results using (i) histogram and (ii) box plot.
- Implement the **Naïve Bayes classification** algorithm using the dataset. Display the classification report with the accuracy.

```
# a) Conduct Exploratory Data Analysis (EDA)
import pandas as pd
df = pd.read_csv('/content/Wine_Quality.csv') # Load the Wine Quality dataset
print(df.head()) # Check the first few rows
print(df.isnull().sum()) # Check for missing values
print(df.describe()) # Summary statistics of the dataset
print(df['quality'].value_counts()) # Distribution of the target variable 'quality'
```

	type	fixed acidity	volatile acidity	citric acid	residual sugar \
0	white	7.0	0.27	0.36	20.7
1	white	6.3	0.30	0.34	1.6
2	white	8.1	0.28	0.40	6.9
3	white	7.2	0.23	0.32	8.5
4	white	7.2	0.23	0.32	8.5

	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH \
0	0.045	45.0	170.0	1.0010	3.00
1	0.049	14.0	132.0	0.9940	3.30
2	0.050	30.0	97.0	0.9951	3.26
3	0.058	47.0	186.0	0.9956	3.19
4	0.058	47.0	186.0	0.9956	3.19

	sulphates	alcohol	quality
0	0.45	8.8	6
1	0.49	9.5	6
2	0.44	10.1	6
3	0.40	9.9	6
4	0.40	9.9	6

	type	0
fixed acidity		10
volatile acidity		8
citric acid		3
residual sugar		2

```

chlorides      2
free sulfur dioxide  0
total sulfur dioxide  0
density        0
pH             9
sulphates      4
alcohol        0
quality        0
dtype: int64
fixed acidity  volatile acidity  citric acid  residual sugar \
count  6487.000000    6489.000000  6494.000000   6495.000000
mean    7.216579      0.339691   0.318722     5.444326
std     1.296750      0.164649   0.145265     4.758125
min     3.800000      0.080000   0.000000     0.600000
25%     6.400000      0.230000   0.250000     1.800000
50%     7.000000      0.290000   0.310000     3.000000
75%     7.700000      0.400000   0.390000     8.100000
max    15.900000      1.580000   1.660000    65.800000

chlorides  free sulfur dioxide  total sulfur dioxide   density \
count  6495.000000    6497.000000    6497.000000  6497.000000
mean   0.056042     30.525319     115.744574   0.994697
std    0.035036    17.749400     56.521855   0.002999
min    0.009000     1.000000     6.000000   0.987110
25%    0.038000    17.000000     77.000000   0.992340
50%    0.047000    29.000000    118.000000   0.994890
75%    0.065000    41.000000    156.000000   0.996990
max    0.611000   289.000000    440.000000   1.038980

pH  sulphates  alcohol  quality
count  6488.000000  6493.000000  6497.000000  6497.000000
mean    3.218395   0.531215   10.491801   5.818378
std     0.160748   0.148814   1.192712   0.873255
min     2.720000   0.220000   8.000000   3.000000
25%     3.110000   0.430000   9.500000   5.000000
50%     3.210000   0.510000  10.300000   6.000000
75%     3.320000   0.600000  11.300000   6.000000
max     4.010000   2.000000  14.900000   9.000000
quality
6  2836
5  2138
7  1079
4  216
8  193
3  30
9  5
Name: count, dtype: int64
addCode
addText

```

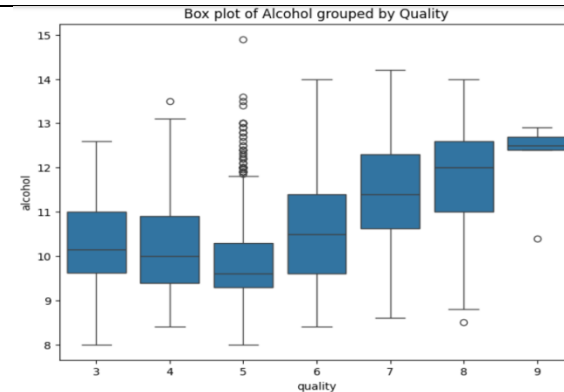
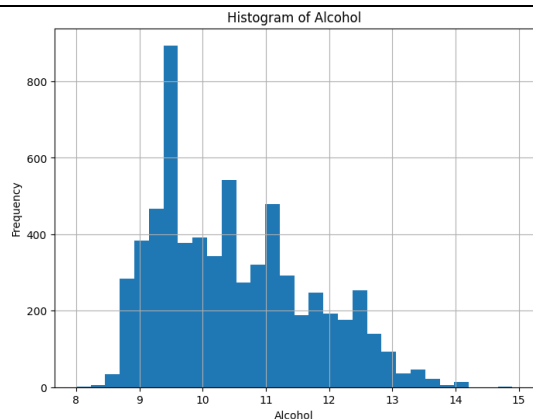
#b) Visualizations

```

import matplotlib.pyplot as plt
import seaborn as sns
# (i) Histogram for a feature like 'alcohol'
plt.figure(figsize=(8, 6))
df['alcohol'].hist(bins=30)
plt.title("Histogram of Alcohol")
plt.xlabel("Alcohol")

```

```
plt.ylabel("Frequency")
plt.show()
# (ii) Box plot for 'alcohol' grouped by 'quality'
plt.figure(figsize=(8, 6))
sns.boxplot(x='quality', y='alcohol', data=df)
plt.title("Box plot of Alcohol grouped by Quality")
plt.show()
```



c) Implement Naïve Bayes Classification

```
from sklearn.model_selection import train_test_split
from sklearn.naive_bayes import GaussianNB
from sklearn.metrics import classification_report, accuracy_score
from sklearn.impute import SimpleImputer # import the imputer
# Step 1: Convert 'quality' into binary class (e.g., good quality = 1 if quality >= 7, otherwise bad quality = 0)
df['quality_label'] = df['quality'].apply(lambda x: 1 if x >= 7 else 0)
# Step 2: Prepare features (X) and target (y)
# drop 'type', 'quality', and the newly created 'quality_label' for the features
X = df.drop(columns=['type', 'quality', 'quality_label'])
y = df['quality_label']
# Step 3: Split the dataset into training and test sets
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2,
random_state=42)
# Step 4: Impute missing values using SimpleImputer
imputer = SimpleImputer(strategy='mean') # create an imputer object with strategy 'mean'
X_train = imputer.fit_transform(X_train) # fit and transform on the training data
X_test = imputer.transform(X_test) # transform the test data
# Step 5: Train the Naive Bayes classifier
nb_model = GaussianNB()
nb_model.fit(X_train, y_train)
# Step 6: Make predictions on the test set
y_pred = nb_model.predict(X_test)
# Step 7: Evaluate the model
accuracy = accuracy_score(y_test, y_pred)
classification_rep = classification_report(y_test, y_pred)
# Output the results
print("Accuracy: {:.2f}%".format(accuracy * 100))
```

```
print("\nClassification Report:\n", classification_rep)
```

```
Accuracy: 77.85%
```

```
Classification Report:
      precision    recall  f1-score   support

     0       0.91      0.81      0.85      1047
     1       0.45      0.66      0.54       253

 accuracy: 0.78      1300
 macro avg: 0.68      1300
 weighted avg: 0.82      1300
```

Program 15: Apply the **Decision Tree Classifier** on the **Wine quality (winequality-red.csv)** dataset to generate the following results.

- (i) Classification Report
- (ii) Confusion Matrix
- (iii) Plot Decision Tree

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

```
# Load the Winequality dataset
```

```
wine_data = pd.read_csv("/content/winequality-red.csv", sep=';')
```

```
print(wine_data.head()) # Check the first five rows
```

```
fixed acidity  volatile acidity  citric acid  residual sugar  chlorides
0           7.4             0.70         0.00           1.9         0.076
1           7.8             0.88         0.00           2.6         0.098
2           7.8             0.76         0.04           2.3         0.092
3          11.2             0.28         0.56           1.9         0.075
4           7.4             0.70         0.00           1.9         0.076

free sulfur dioxide  total sulfur dioxide  density  pH  sulphates \
0             11.0             34.0    0.9978  3.51      0.56
1             25.0             67.0    0.9968  3.20      0.68
2             15.0             54.0    0.9970  3.26      0.65
3             17.0             60.0    0.9980  3.16      0.58
4             11.0             34.0    0.9978  3.51      0.56

alcohol  quality
0       9.4       5
1       9.8       5
2       9.8       5
3       9.8       6
4       9.4       5
```

```
# Check if the 'color' column exists in the dataset
```

```
if 'color' in wine_data.columns:
```

```
    # Convert the 'color' column to numerical values using one-hot encoding
```

```
wine_data = pd.get_dummies(wine_data, columns=['color'], drop_first=True)
```

```
# Separate features (X) and target variable (y)
```

```
X = wine_data.drop('quality', axis=1)
```

```
y = wine_data['quality']
```

```
# Split the dataset into training and testing sets
```

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

```
# Initialize the Decision Tree Classifier with max_depth to prune the tree
```

```
dt_classifier = DecisionTreeClassifier(random_state=42, max_depth=3)
```

```
dt_classifier.fit(X_train, y_train) # Train the model
```

DecisionTreeClassifier
DecisionTreeClassifier(criterion='entropy', max_depth=3, random_state=42)

```

# (i) Classification Report
y_pred = dt_classifier.predict(X_test)
classification_rep = classification_report(y_test, y_pred, zero_division=1)
print("Classification Report:")
print(classification_rep)

```

```

Classification Report:
              precision    recall  f1-score   support

     3           1.00        0.00        0.00         1
     4           1.00        0.00        0.00        10
     5           0.54        0.91        0.68       130
     6           0.53        0.31        0.39       132
     7           0.42        0.24        0.30        42
     8           1.00        0.00        0.00         5

 accuracy          0.53        0.53        0.53       320
 macro avg          0.75        0.24        0.23       320
 weighted avg          0.54        0.53        0.48       320

```

```

# (ii) Confusion Matrix
conf_matrix = confusion_matrix(y_test, y_pred)
print("\nConfusion Matrix:")
print(conf_matrix)

```

```

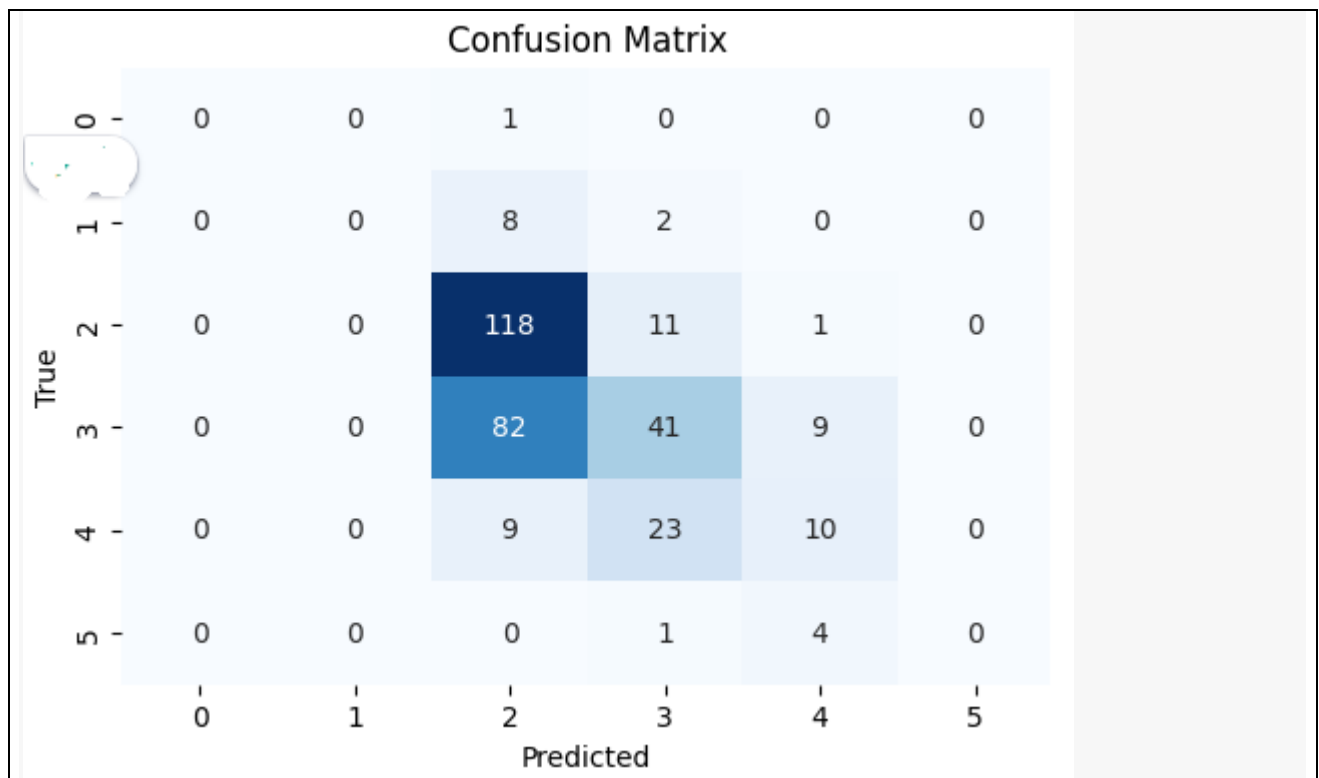
Confusion Matrix:
[[ 0  0  1  0  0  0]
 [ 0  0  8  2  0  0]
 [ 0  0 118 11  1  0]
 [ 0  0  82 41  9  0]
 [ 0  0   9 23 10  0]
 [ 0  0   0  1  4  0]]

```

```

# Visualize the Confusion Matrix using a heatmap
plt.figure(figsize=(6, 4))
sns.heatmap(conf_matrix, annot=True, fmt='d', cmap='Blues', cbar=False)
plt.xlabel('Predicted')
plt.ylabel('True')
plt.title('Confusion Matrix')
plt.show()

```

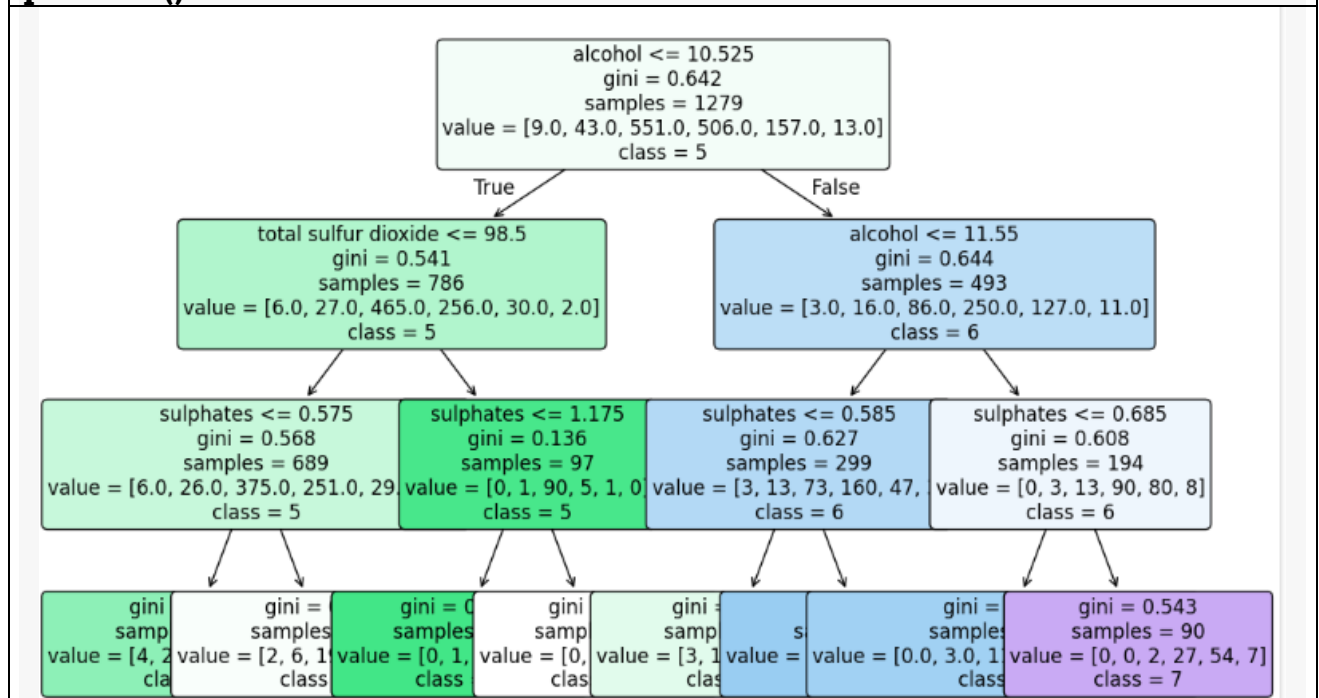



(iii) Plot Pruned Decision Tree

plt.figure(figsize=(12, 8))

tree.plot_tree(dt_classifier, feature_names=X.columns, class_names=[str(label) for label in dt_classifier.classes_], filled=True, rounded=True)

plt.show()



Program 16: Apply the **Decision Tree Classifier** on **iris dataset** to generate the following results.

- (i) Classification Report
- (ii) Confusion Matrix
- (iii) Plot Decision Tree

```
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 classification_report, confusion_matrix
import matplotlib.pyplot as plt
import seaborn as sns

iris = load_iris() # Load the Iris dataset
X = pd.DataFrame(iris.data, columns=iris.feature_names) # Features
y = pd.Series(iris.target) # Target variable
# Split the dataset into training and test sets (80% train, 20% test)
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2,
random_state=42)

# Initialize the Decision Tree Classifier
clf = DecisionTreeClassifier (criterion='entropy', max_depth=3,
random_state=42)
clf.fit(X_train, y_train) # Train the classifier on the training data
y_pred = clf.predict(X_test) # Make predictions on the test data

# (i) Classification Report
print("Classification Report:")
print(classification_report(y_test, y_pred))

Classification Report:
              precision    recall  f1-score   support

     0         1.00      1.00      1.00        10
     1         1.00      1.00      1.00         9
     2         1.00      1.00      1.00        11

 accuracy          1.00          1.00          1.00        30
 macro avg          1.00          1.00          1.00        30
weighted avg          1.00          1.00          1.00        30

# (ii) Confusion Matrix
cm = confusion_matrix(y_test, y_pred)
print("Confusion Matrix:")
print(cm)
```

Confusion Matrix:

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

Visualize the Confusion Matrix using a heatmap

```
plt.figure(figsize=(6, 4))
```

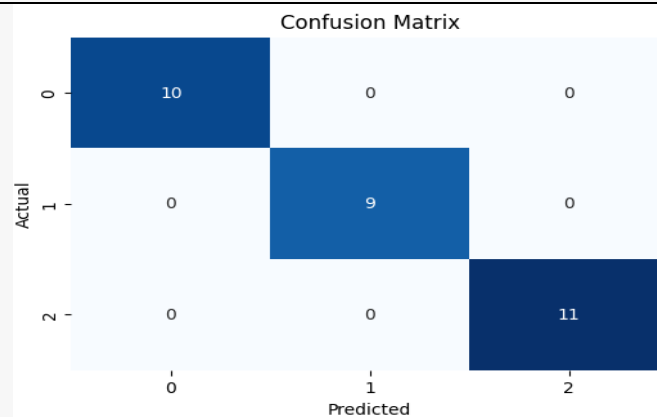
```
sns.heatmap(cm, annot=True, fmt='d', cmap='Blues', cbar=False)
```

```
plt.xlabel('Predicted')
```

```
plt.ylabel('True')
```

```
plt.title('Confusion Matrix')
```

```
plt.show()
```



(iii) Plot the Decision Tree

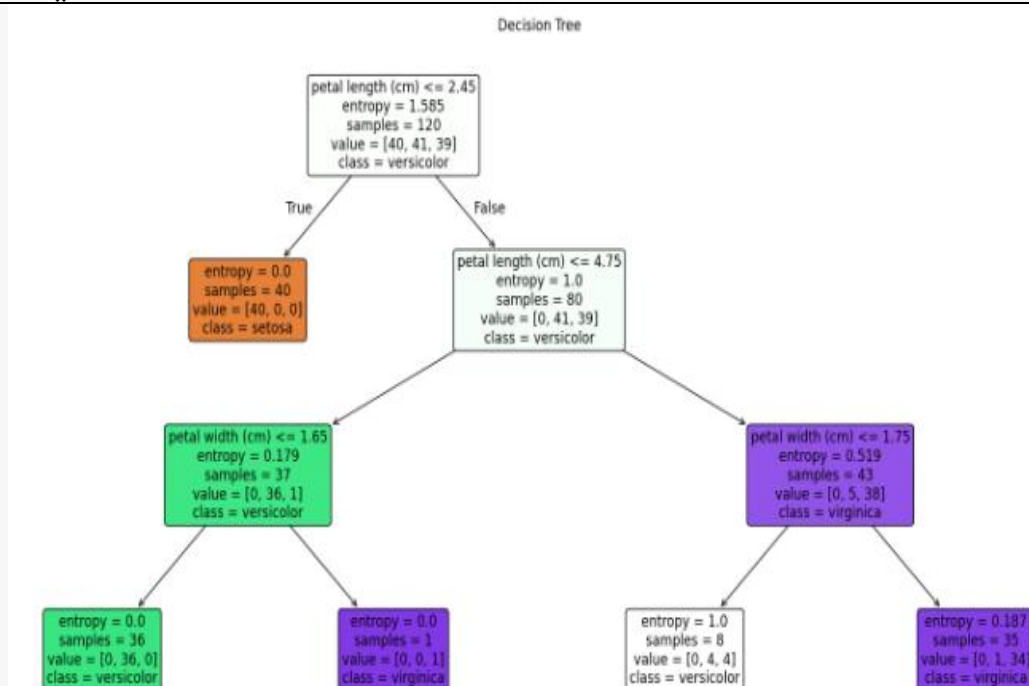
```
plt.figure(figsize=(20, 10))
```

```
plot_tree(clf, filled=True, feature_names=iris.feature_names,
```

```
class_names=iris.target_names, rounded=True, fontsize=12)
```

```
plt.title('Decision Tree')
```

```
plt.show()
```



Program 17: Implement Simple linear regression for the data sets 'student_score.csv'

- **Linear Regression (Simple Linear Regression):** Only 1 independent variable & 1 dependent variable.

```
# Import necessary libraries
import numpy as np #Provides support for numerical operations, particularly arrays.
import pandas as pd #for handling and analyzing data in DataFrames, making it easy to load, manipulate, and process datasets.
import matplotlib.pyplot as plt
from sklearn.model_selection import train_test_split
from sklearn.linear_model import LinearRegression
from sklearn.metrics import mean_squared_error, r2_score
# LinearRegression: A class in sklearn.linear_model that enables fitting a linear model to data.
#mean_squared_error, r2_score: Functions in sklearn.metrics used to evaluate model performance.
```

```
# Loads the student_scores.csv dataset into a pandas DataFrame named student.
student = pd.read_csv('/content/student_scores.csv')
# Display the first few rows and basic info about the dataset
print(student.head()) #Displays the first 5 rows of the dataset
print(student.describe()) #Provides summary statistics (like mean, min, max, etc.)
print(student.info()) #information about each column, including the data type and non-null count
```

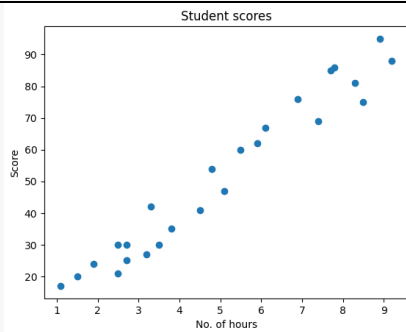
```
Hours Scores
0 2.5 21
1 5.1 47
2 3.2 27
3 8.5 75
4 3.5 30

Hours Scores
count 25.000000 25.000000
mean 5.012000 51.480000
std 2.525094 25.286887
min 1.100000 17.000000
25% 2.700000 30.000000
50% 4.800000 47.000000
75% 7.400000 75.000000
max 9.200000 95.000000
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 25 entries, 0 to 24
Data columns (total 2 columns):
# Column Non-Null Count Dtype
---
0 Hours 25 non-null float64
1 Scores 25 non-null int64
dtypes: float64(1), int64(1)
memory usage: 528.0 bytes
None
```

```
# Separate the features (No. of hours) and target (Score)
Xax = student.iloc[:, 0] # Selects the first column (No. of hours) as the independent variable (Xax).
Yax = student.iloc[:, 1] # Selects the second column (Score) as the dependent variable (Yax).

# Visualize the data with a scatter plot
plt.scatter(Xax, Yax)
plt.xlabel("No. of hours")
plt.ylabel("Score")
```

```
plt.title("Student scores")
plt.show()
```



```
# Prepare data for training and testing
x = student.iloc[:, :-1] #Selects all columns except the last one as the independent variable (hours studied).
y = student.iloc[:, 1]    # Selects the last column as the dependent variable (score).
print('x values :\n', x)
print('y values :\n', y)
```

```
x values :
```

```
Hours
```

```
0    2.5
1    5.1
2    3.2
3    8.5
4    3.5
5    1.5
6    9.2
7    5.5
8    8.3
9    2.7
10   7.7
11   5.9
12   4.5
13   3.3
14   1.1
15   8.9
16   2.5
17   1.9
18   6.1
19   7.4
20   2.7
21   4.8
22   3.8
23   6.9
24   7.8
```

```
y values :
```

```
0    21
1    47
2    27
3    75
4    30
5    20
6    88
7    60
8    81
9    25
10   85
```

```

11    62
12    41
13    42
14    17
15    95
16    30
17    24
18    67
19    69
20    30
21    54
22    35
23    76
24    86

```

```
Name: Scores, dtype: int64
```

```

# Split the dataset into training and testing sets
x_train, x_test, y_train, y_test = train_test_split(x, y, test_size=0.2,
random_state=42)

# Train the Linear Regression model
regressor = LinearRegression() #Initializes the linear regression model.
regressor.fit(x_train, y_train) # Trains (fits) the model on the training data, finding the best- # fit line
that predicts scores based on hours studied.
# Output the intercept and coefficient
print('INTERCEPT = ', regressor.intercept_) #Retrieves the y-intercept of the regression line.
print('COEFFICIENT = ', regressor.coef_[0]) #Retrieves the slope (coefficient) of the regression line
#These values represent the linear equation: Score = (Coefficient * Hours) + Intercept

```

```
INTERCEPT = 2.826892353899737
```

```
COEFFICIENT = 9.682078154455697
```

```

# Make predictions on the test set
y_pred = regressor.predict(x_test) #Uses the trained model to predict scores (y_pred) for
# the test set based on hours studied in x_test.
# Print the actual vs predicted values
print("Actual vs Predicted values:")
for actual, predicted in zip(y_test, y_pred):
    print("Actual value:", actual, "Predicted value:", predicted)
#Iterates over each actual and predicted value pair, printing them for comparison.

```

```
Actual vs Predicted values:
```

```
Actual value: 81 Predicted value: 83.18814103588203
```

```
Actual value: 30 Predicted value: 27.03208774003898
```

```
Actual value: 21 Predicted value: 27.03208774003898
```

```
Actual value: 76 Predicted value: 69.63323161964405
```

```
Actual value: 62 Predicted value: 59.951153465188355
```

```

# Calculate and print the number of mislabeled points
mislabeled_points = np.sum(np.round(y_test) != np.round(y_pred))
#Counts the number of mismatched points between the actual (y_test) and predicted (y_pred) values
#(after rounding to avoid small numerical differences).
print("Number of mislabeled points from test data set:", mislabeled_points)

```

```
Number of mislabeled points from test data set: 5
```

```

# Evaluate the model performance
mse = mean_squared_error(y_test, y_pred) #Calculates the Mean Squared Error (MSE) between #actual

```

and predicted values, a common metric for regression that measures the average squared difference between #predicted and actual values.

```
r2 = r2_score(y_test, y_pred) # Calculates the R-squared score, which indicates how well the model fits #the data (values closer to 1 indicate a better fit).
```

```
print("Mean Squared Error:", mse)
```

```
print("R-squared:", r2)
```

```
Mean Squared Error: 18.943211722315272
```

```
R-squared: 0.9678055545167994
```

Program 18: Implement **multiple linear regression** for the data sets "Company_data.csv"

- **Two or more independent variables** to predict the dependent variable.

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
from sklearn.model_selection import train_test_split
from sklearn.linear_model import LinearRegression
from sklearn import metrics
import pandas as pd

# Load the dataset
advertising = pd.read_csv('/content/Advertising.csv')
# Display the first few rows and basic info about the dataset
print(advertising.head())
print(advertising.describe())
print(advertising.info())
```

	TV	Radio	Newspaper	Sales
0	230.1	37.8	69.2	22.1
1	44.5	39.3	45.1	10.4
2	17.2	45.9	69.3	12.0
3	151.5	41.3	58.5	16.5
4	180.8	10.8	58.4	17.9

```
count 200.000000 200.000000 200.000000 200.000000
mean 147.042500 23.264000 30.554000 15.130500
std 85.854236 14.846809 21.778621 5.283892
min 0.700000 0.000000 0.300000 1.600000
25% 74.375000 9.975000 12.750000 11.000000
50% 149.750000 22.900000 25.750000 16.000000
75% 218.825000 36.525000 45.100000 19.050000
max 296.400000 49.600000 114.000000 27.000000
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 200 entries, 0 to 199
Data columns (total 4 columns):
# Column Non-Null Count Dtype
---
0 TV 200 non-null float64
1 Radio 200 non-null float64
2 Newspaper 200 non-null float64
3 Sales 200 non-null float64
dtypes: float64(4)
memory usage: 6.4 KB
None
addCode
addText
```

```
# Separate the features (independent variables) and the target variable (dependent variable)
print("Feature values:")
x = advertising.iloc[:, :-1] # Selects all columns except the last one as features
print(x)
```

	TV	Radio	Newspaper
0	230.1	37.8	69.2
1	44.5	39.3	45.1


```

2      17.2    45.9    69.3
3     151.5    41.3    58.5
4     180.8    10.8    58.4
..      ...      ...      ...
195    38.2     3.7    13.8
196    94.2     4.9     8.1
197   177.0     9.3     6.4
198   283.6    42.0    66.2
199   232.1     8.6     8.7

```

[200 rows x 3 columns]

```

print("Target variable values:")
y = advertising.iloc[:, -1]  # Selects the last column as the target
variable
print(y)

```

Target variable values:

```

0      22.1
1      10.4
2      12.0
3      16.5
4      17.9
...
195     7.6
196    14.0
197    14.8
198    25.5
199    18.4

```

Name: Sales, Length: 200, dtype: float64

```

# Split the dataset into training and testing sets (70% training, 30%
testing)
x_train, x_test, y_train, y_test = train_test_split(x, y,
test_size=0.3, random_state=42)

```

```

# Train the Multiple Linear Regression model
regressor = LinearRegression()
regressor.fit(x_train, y_train)
# Output the intercept and coefficients of the regression model
print("Intercept:", regressor.intercept_)
print("Coefficients:", regressor.coef_)

```

Intercept: 4.743766701589685

Coefficients: [0.05358869 0.10270677 0.00793167]

```

# Make predictions on the test set
y_pred = regressor.predict(x_test)
# Print actual vs predicted values for the test set
print("Actual vs Predicted values:")
for actual, predicted in zip(y_test, y_pred):
    print("Actual value:", actual, "Predicted value:", predicted)

```

Actual value: 21.4 Predicted value: 23.689143963340577

Actual value: 7.3 Predicted value: 9.519145500497697

Actual value: 24.7 Predicted value: 21.607368359108364

Actual value: 12.6 Predicted value: 12.781013179417098

Actual value: 22.3 Predicted value: 21.086363448001876

Actual value: 8.4 Predicted value: 8.760542459543231

```

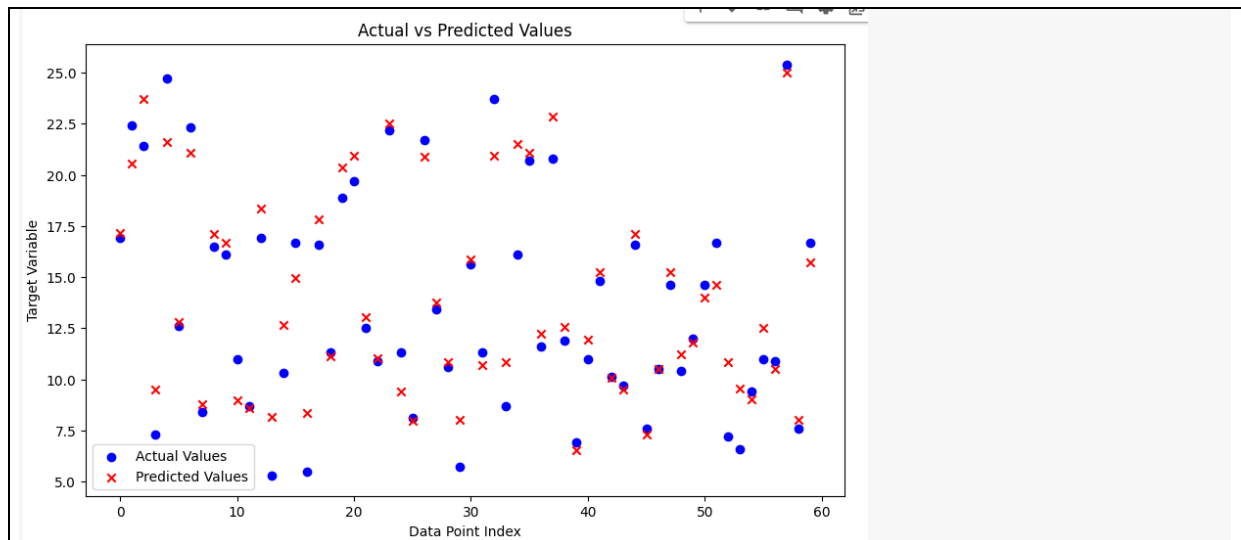
Actual value: 16.5 Predicted value: 17.11499950978093
Actual value: 16.1 Predicted value: 16.687896360018495
Actual value: 11.0 Predicted value: 8.975846634860197
----
---
Actual value: 6.6 Predicted value: 9.55839414535506
Actual value: 9.4 Predicted value: 9.03749681483222
Actual value: 11.0 Predicted value: 12.511833134002398
Actual value: 10.9 Predicted value: 10.525510210425061
Actual value: 25.4 Predicted value: 25.019008241540465
Actual value: 7.6 Predicted value: 7.993349434647646
Actual value: 16.7 Predicted value: 15.73916263437171

# Calculate and print the number of mislabeled points
mislabeled_points = np.sum(np.round(y_test) != np.round(y_pred))
print("Number of mislabeled points from test data set:", mislabeled_points)
Number of mislabeled points from test data set: 37

# Calculate and print evaluation metrics
mae = metrics.mean_absolute_error(y_test, y_pred)
mse = metrics.mean_squared_error(y_test, y_pred)
rmse = np.sqrt(mse)
print("Mean Absolute Error:", mae)
print("Mean Squared Error:", mse)
print("Root Mean Squared Error:", rmse)
Mean Absolute Error: 1.1594875061090582
Mean Squared Error: 2.541624036229147
Root Mean Squared Error: 1.5942471691143587

# Visualization of Actual vs Predicted values
plt.figure(figsize=(10, 6))
plt.scatter(range(len(y_test)), y_test, color='blue', label='Actual
Values')
plt.scatter(range(len(y_pred)), y_pred, color='red', label='Predicted
Values', marker='x')
plt.title('Actual vs Predicted Values')
plt.xlabel('Data Point Index')
plt.ylabel('Target Variable')
plt.legend()
plt.show()

```



Program 19: Given dataset contains 200 records and five columns, two of which describe the customer's annual income and spending score. The latter is a value from 0 to 100. The higher the number, the more this customer has spent with the company in the past: Using k means clustering creates 6 clusters of customers based on their spending pattern.

- Visualize the same in a scatter plot with each cluster in a different color scheme.
- Display the cluster labels of each point.(print cluster indexes)
- Display the cluster centers.

```
# Import necessary libraries
import pandas as pd
import matplotlib.pyplot as plt
from sklearn.cluster import KMeans
import numpy as np
# Load customer data
customer = pd.read_csv('/content/Customer_Data.csv')

# Extract the 'Annual Income' and 'Spending Score' columns for clustering
points = customer.iloc[:, 3:5].values # Assuming columns 3 and 4 are 'Annual Income' and 'Spending Score'
x = points[:, 0]
y = points[:, 1]
# Plot initial data points
plt.figure(figsize=(8, 6))
plt.scatter(x, y, s=50, alpha=0.7, c='blue')
plt.xlabel('Annual Income (k$)')
plt.ylabel('Spending Score (1-100)')
plt.title('Customer Data: Annual Income vs Spending Score')
plt.show()
# Function to apply KMeans clustering and plot results
def kmeans_clustering_and_plot(points, n_clusters):
    # Apply KMeans
    kmeans = KMeans(n_clusters=n_clusters, random_state=0)
```

```

kmeans.fit(points)
cluster_labels = kmeans.predict(points)
cluster_centers = kmeans.cluster_centers_

# Plot clustered data
plt.figure(figsize=(8, 6))
scatter = plt.scatter(x, y, c=cluster_labels, s=50, alpha=0.7,
cmap='viridis')
plt.scatter(cluster_centers[:, 0], cluster_centers[:, 1], s=200, c='red',
marker='X', label='Centroids')
plt.xlabel('Annual Income (k$)')
plt.ylabel('Spending Score (1-100)')
plt.title(f'K-Means Clustering with {n_clusters} Clusters')
plt.legend()
plt.colorbar(scatter, label="Cluster Label")
plt.show()
# Print cluster labels for each point
print(f'Cluster labels for K={n_clusters}:\n', cluster_labels)
print(f'Cluster centers for K={n_clusters}:\n', cluster_centers)
# Visualize clustering for different values of K
for k in range(4, 9): # Testing for K=4 to K=8
    kmeans_clustering_and_plot(points, n_clusters=k)

```

Program 20: Write a program to implement a simple web crawler using Python. Extract and display the content of the page(p tag) .

- A **web crawler**, also known as a **spider** or **bot**, is an automated program that systematically browses the web to collect information from websites.

```

import requests
from bs4 import BeautifulSoup

print("Batch : MCA 2023-25")

# Function to get data from the URL
def getdata(url):
    r = requests.get(url)
    return r.content

# URL to crawl
url="https://www.w3schools.com/python/default.asp"
htmldata = getdata(url)

# Parse the HTML data with BeautifulSoup
soup = BeautifulSoup(htmldata, 'html.parser')

# Find all <p> tags and display their content

```

```

paragraphs = soup.find_all('p')
print("<P> tag count:", len(paragraphs)) # Display the count of <p> tags

for p in paragraphs:
    print(p.get_text()) # Print the text content of each <p> tag

```

Program 21 : Implement the **K-Means clustering** algorithm using the <Iris.csv> dataset

- a) Conduct exploratory data analysis on the given dataset and report the details.
- b) Visualize the analysis results using (i) scatter plot (ii) histogram & (iii) box plot.
- c) Try with different K values and plot the elbow graph for the k values.

Program 22: Dataset: <Housing_Price.csv>

- a) Conduct exploratory data analysis on the given dataset and report the details.
- b) Visualize the analysis results using (i) scatter plot (ii) histogram & (iii) box plot.
- c) Implement the **K-Means clustering** algorithm using the dataset. Try with different k values and plot the elbow graph for the k values.
