

MAHA BARATHI ENGINEERING COLLEGE

NH-79, SALEM-CHENNAI HIGHWAY, A.VASUDEVANUR, CHINNASALEM TK, KALLAKURICHI DT – 606 201.

Approved by AICTE, New Delhi & Affiliated to Anna University, Chennai

2(f) & 12(B) status of UGC, New Delhi,

www.mbec.ac.in | 04151-256333, 257333 | mbec123@gmail.com



DEPARTMENT OF COMPUTER SCIENCE AND ENGINEERING

CS3361 – DATA SCIENCE LAB MANUAL

II Year/III Semester B.E CSE

Regulation 2021
(As Per Anna University, Chennai syllabus)

Prepared By,

P.AKILA
(AP/CSE)

Verified By,

N.KHADIRKUMAR
(HOD/CSE)

OBJECTIVES:

- To understand the python libraries for data science
- To understand the basic Statistical and Probability measures for data science.
- To learn descriptive analytics on the benchmark data sets.
- To apply correlation and regression analytics on standard data sets.
- To present and interpret data using visualization packages in Python .

LIST OF EXPERIMENTS:

1. Download, install and explore the features of NumPy, SciPy, Jupyter, Statsmodels and Pandas packages.
2. Working with Numpy arrays
3. Working with Pandas data frames
4. Reading data from text files, Excel and the web and exploring various commands for doing descriptive analytics on the Iris data set.
5. Use the diabetes data set from UCI and Pima Indians Diabetes data set for performing the following:
 - a. Univariate analysis: Frequency, Mean, Median, Mode, Variance, Standard Deviation, Skewness and Kurtosis.
 - b. Bivariate analysis: Linear and logistic regression modeling
 - c. Multiple Regression analysis
 - d. Also compare the results of the above analysis for the two data sets.
6. Apply and explore various plotting functions on UCI data sets.
 - a. Normal curves
 - b. Density and contour plots
 - c. Correlation and scatter plots
 - d. Histograms
 - e. Three dimensional plotting
7. Visualizing Geographic Data with Basemap

TOTAL : 60 PERIODS**OUTCOMES:****On completion of this course, the students will be able to:****CO1:** Make use of the python libraries for data science**CO2:** Make use of the basic Statistical and Probability measures for data science.**CO3:** Perform descriptive analytics on the benchmark data sets.**CO4:** Perform correlation and regression analytics on standard data sets**CO5:** Present and interpret data using visualization packages in Python.

INSTALLING ANACONDA ON WINDOWS

Anaconda distribution of Python is the best option for problem solvers who want to use Python. Anaconda is free (although the download is large which can take time) and can be installed. Anaconda comes bundled with about 600 packages pre-installed including NumPy, Matplotlib and SymPy. These three packages are very useful for problem solvers and will be discussed in subsequent chapters.

Follow the steps below to install the Anaconda distribution of Python on Windows.

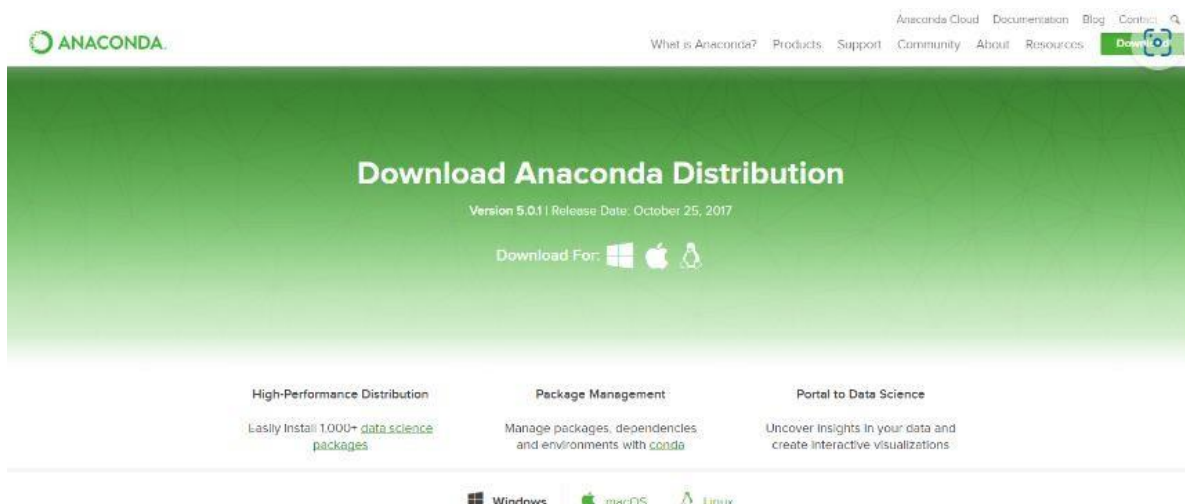
Steps:

1. Visit [Anaconda.com/downloads](https://anaconda.com/downloads)
2. Select Windows
3. Download the .exe installer
4. Open and run the .exe installer
5. Open the Anaconda Prompt and run some Python code

1. Visit the Anaconda downloads page

Go to the following link: [Anaconda.com/downloads](https://anaconda.com/downloads)

The Anaconda Downloads Page will look something like this:



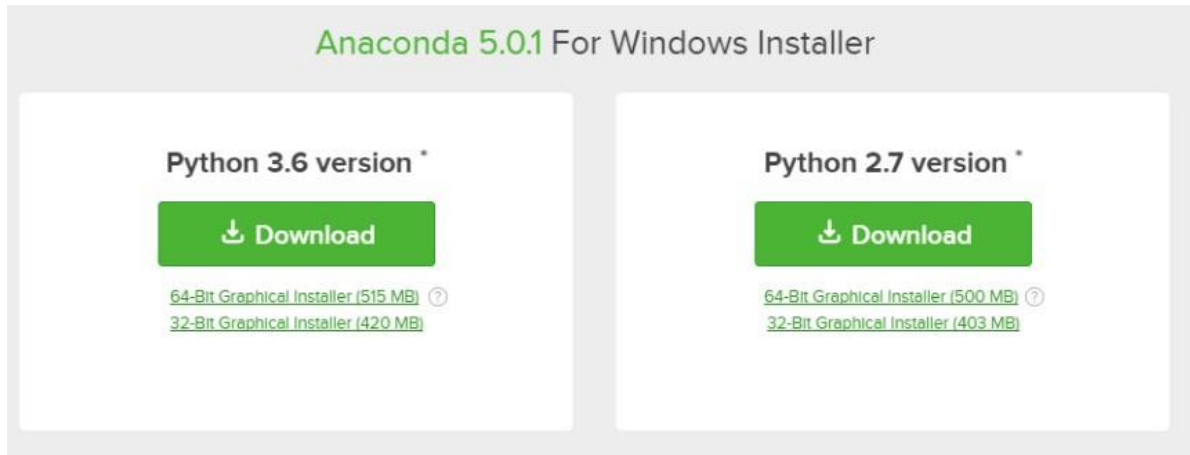
2. Select Windows

Select Windows where the three operating systems are listed.

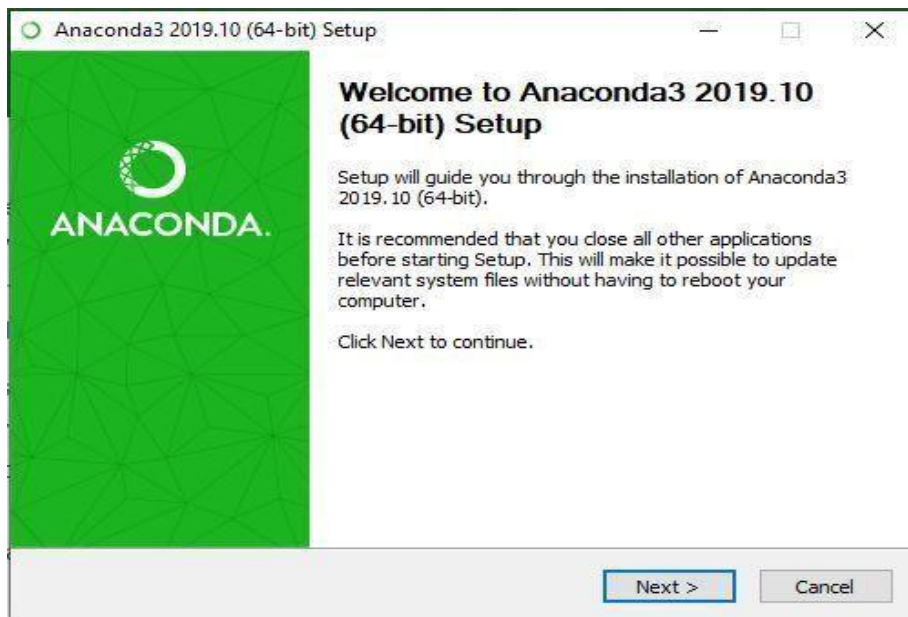


3.Download

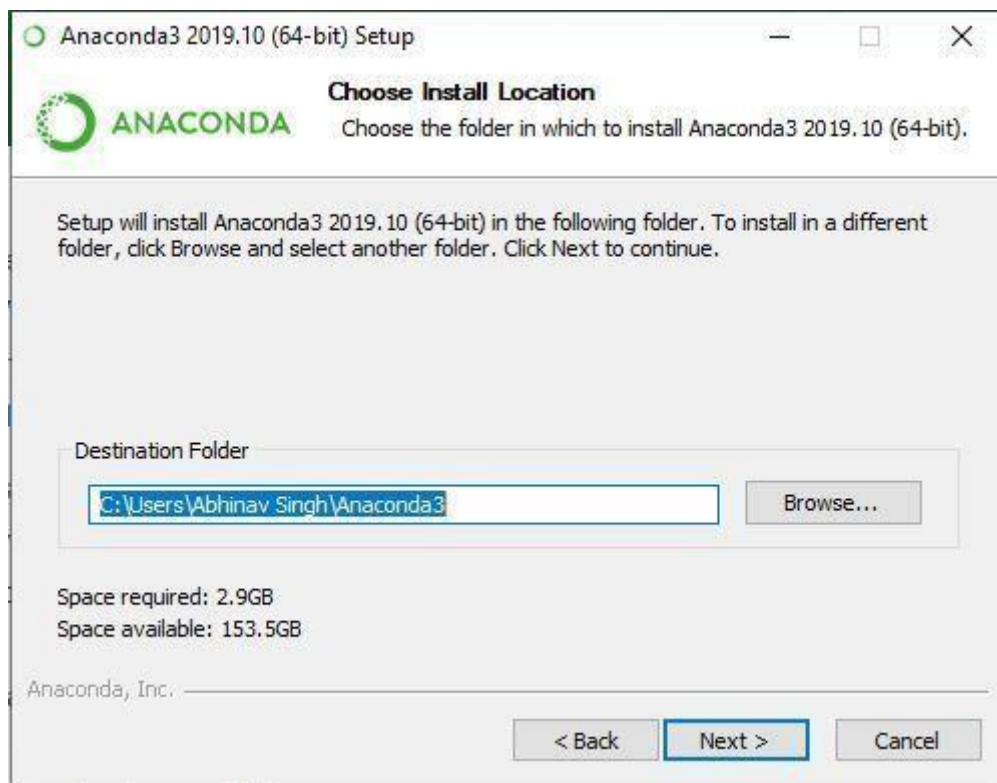
Download the most recent Python 3 release. At the time of writing, the most recent release was the Python Version. Python 2.7 is legacy Python. For problem solvers, select the Python 3.6 version. If you are unsure if your computer is running a 64-bit or 32-bit version of Windows, select 64-bit as 64-bit Windows is most common.



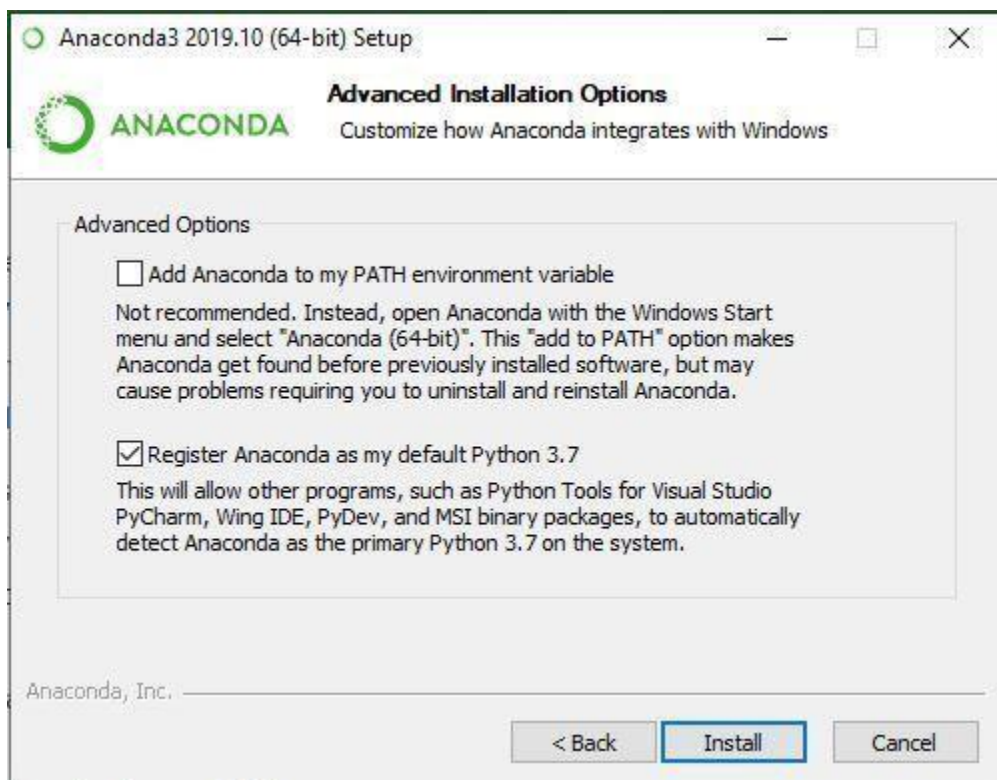
Begin with the installation process:
Getting Started:



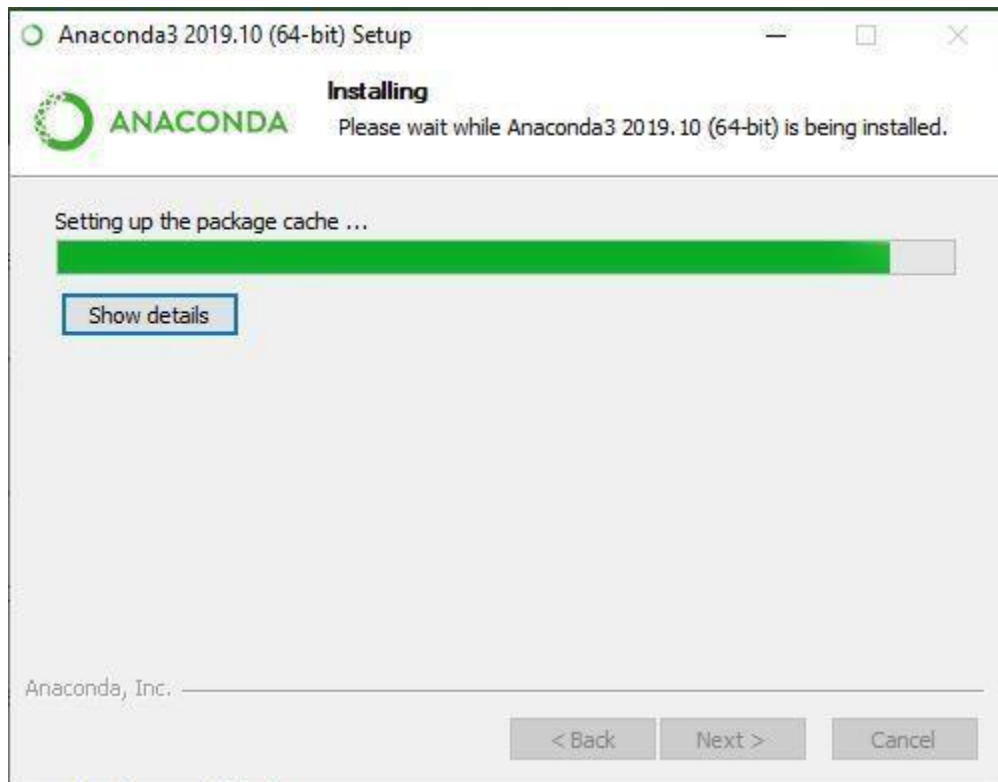
Choose Installation Location:



- **Advanced Installation Option:**



- Getting through the Installation Process:



- Recommendation to Install Pycharm:

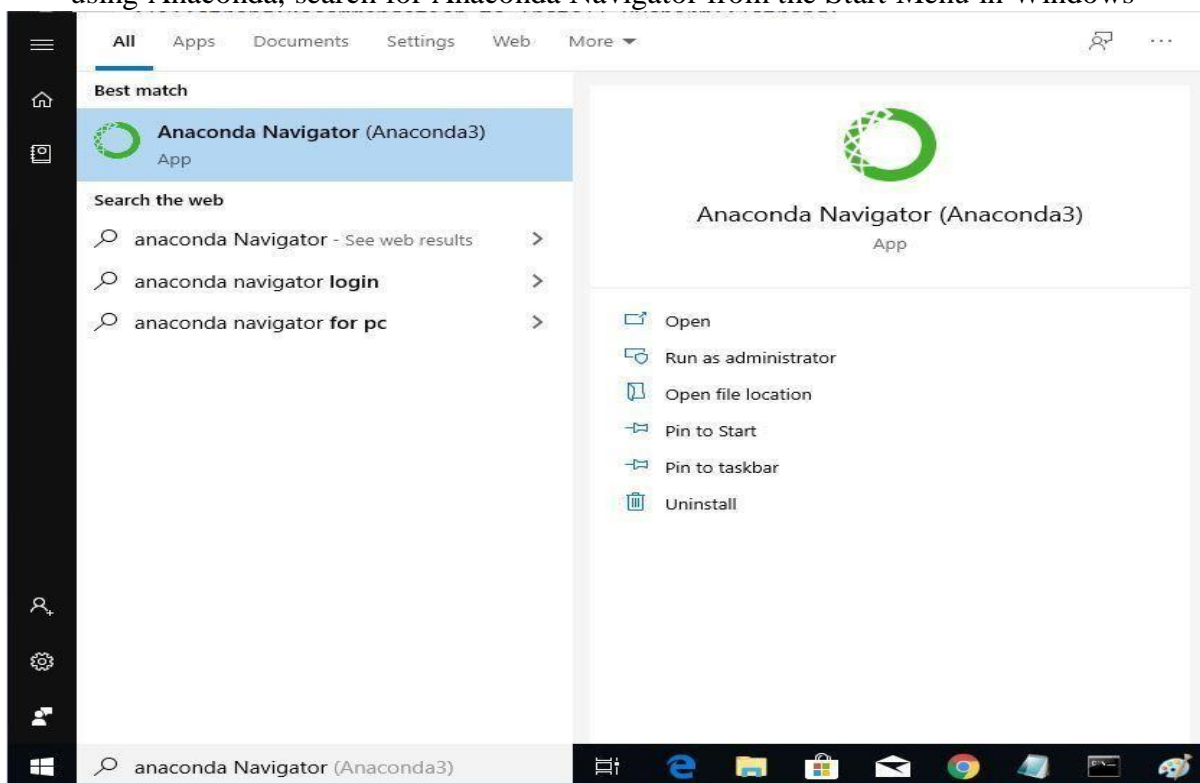


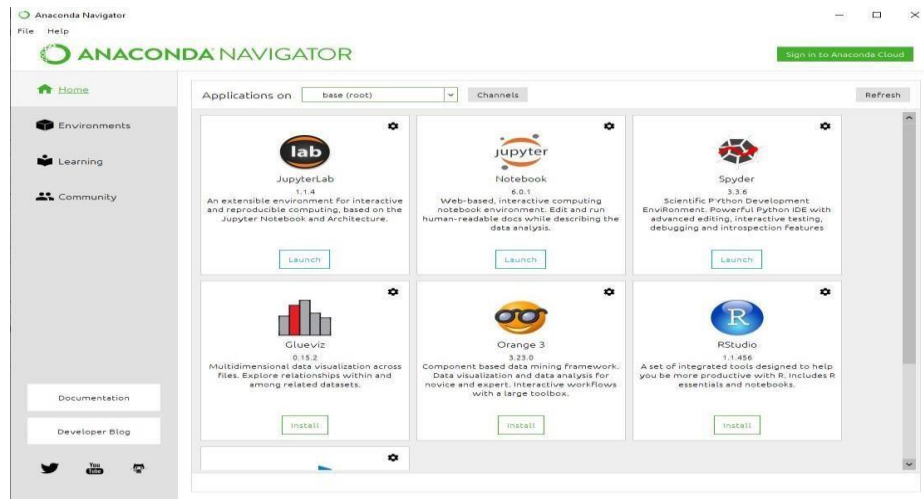
- Finishing up the Installation:



Working with Anaconda:

Once the installation process is done, Anaconda can be used to perform multiple operations. To begin using Anaconda, search for Anaconda Navigator from the Start Menu in Windows





Exploring NumPy Packages:

NumPy is a Python package used for numerical computation. NumPy is one of the foundational packages for scientific computing with Python. NumPy's core data type is the array and NumPy functions operate on arrays.

Installing NumPy

Before NumPy's functions and methods can be used, NumPy must be installed. Depending on which distribution of Python you use, the installation method is slightly different.

Install NumPy on Anaconda

If you installed the Anaconda distribution of Python, NumPy comes pre-installed and no further installation steps are necessary.

If you use a version of Python from python.org or a version of Python that came with your operating system, the Anaconda Prompt and conda or pip can be used to install NumPy.

Install NumPy with the Anaconda Prompt

To install NumPy, open the Anaconda Prompt and type:

```
> conda install numpy
```

Type y for yes when prompted.

Verify NumPy installation

To verify NumPy is installed, invoke NumPy's version using the Python REPL. Import NumPy and call the `version` attribute common to most Python packages.

```
In [1]:
import numpy as np
np.version
```

```
Out[1]:'1.16.4'
```

A version number like '1.16.4' indicates a successful NumPy installation.

Exploring SciPy Packages:

Installing With Pip

You can install SciPy from PyPI with pip:

```
python -m pip install scipy
```

Installing Via Conda

You can install SciPy from the defaults or conda-forge channels with conda:

```
conda install scipy
```

Exploring Jupyter Packages:

Installing Jupyter

The simplest way to install Jupyter notebooks is to download and install the Anaconda distribution of Python. The Anaconda distribution of Python comes with Jupyter notebook included and no further installation steps are necessary.

Installing Jupyter on Windows using the Anaconda Prompt

To install Jupyter on Windows, open the Anaconda Prompt and type:

```
> conda install jupyter
```

Type y for yes when prompted. Once Jupyter is installed, type the command below into the Anaconda Prompt to open the Jupyter notebook file browser and start using Jupyter notebooks.

```
> jupyter notebook
```

Exploring Stats models Packages:

The easiest way to install stats models is to install it as part of the Anaconda distribution, a cross-platform distribution for data analysis and scientific computing. This is the recommended installation method for most users.

Instructions for installing from PyPI, source or a development version are also provided.

Python Support

Stats models supports Python 3.8, 3.9, and 3.10.

Anaconda

Stats models is available through conda provided by Anaconda. The latest release can be installed using:

```
conda install -c conda-forge stats models
```

PyPI (pip)

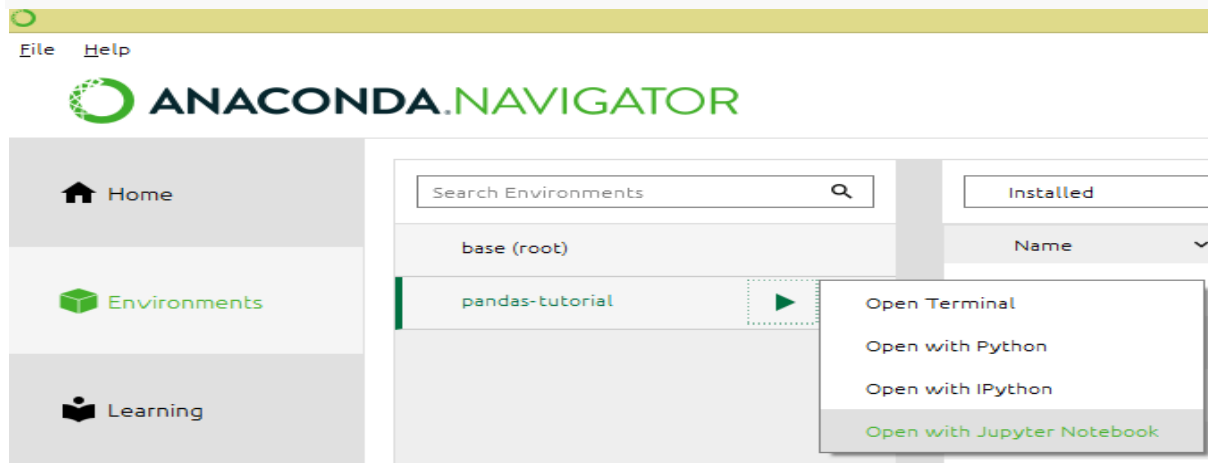
To obtain the latest released version of stats models using pip:

```
python -m pip install stats model.
```

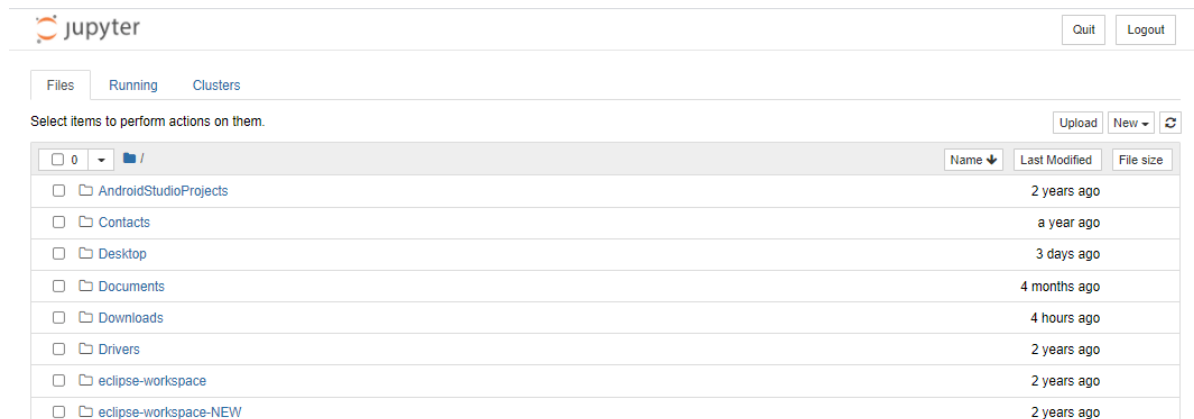
Follow this link to our PyPI page to directly download wheels or source.

Exploring Pandas packages

Go to **Anaconda Navigator -> Environments -> your environment** (mine pandas-tutorial) -> select **Open with Jupyter Notebook**



This opens up Jupyter Notebook in the default browser.



Now select New -> Python X and enter the below lines and select Run.

The screenshot shows a Jupyter Notebook interface. At the top, the title bar reads "jupyter Untitled4 Last Checkpoint: a few seconds ago (unsaved changes)". On the right side of the title bar, there is a "Logout" button and a Python logo. Below the title bar is a menu bar with options: File, Edit, View, Insert, Cell, Kernel, and Help. To the right of the menu bar, it says "Trusted" with a pencil icon and "Python 3 (ipykernel)" with a circular refresh icon. Below the menu bar is a toolbar with icons for file operations (new, open, save, print), navigation (up, down), and execution (run, stop, refresh, next). The main area of the notebook contains three code cells. The first cell has the code `import pandas as pd`. The second cell has the code `pd.__version__`. The output of the second cell is `'1.3.2'`. The third cell is currently empty, showing `In []:` with a cursor.

Result:

This completes installing Anaconda and running pandas on Jupyter Notebook.

EX.NO:2	ARRAY INDEXING using NUMPY
DATE:	

AIM:

To write a python program to implement array indexing using numpy

ALGORITHM:

Step1: Start
Step2: Import necessary libraries-numpy
Step3: Using random module, seed for reproducibility
Step4: Create one dimensional, two dimensional array using randint
Step5: Access the elements by using the index for the different dimensional array. Step6:
Stop the Program

PROGRAM:

```
import numpy as np

np.random.seed(0) # seed for reproducibility

x1 = np.random.randint(10, size=6) # One-dimensional array
x2 = np.random.randint(10, size=(3, 4)) # Two-dimensional array
x3 = np.random.randint(10, size=(3, 4, 5)) # Three-dimensional array
print(x1[0])
print(x1[4])
#To index from the end of the array, negative indices are used
print(x1[-1])
print(x1[-2])

#In a multidimensional array, items are accessed using a comma-separated tuple
```

#of indices:

```
print(x2) print(x2[0,  
0])
```

```
print(x2[2, 0])
```

```
print(x2[2, -1])
```

#modifying values using index notation:

```
x2[0, 0] = 12
```

```
print(x2)
```

```
x1[0] = 3.14159 # this will be truncated!
```

```
print(x1)
```

OUTPUT

```
[5 0 3 3 7 9]
```

```
5
```

```
7
```

```
9
```

```
7
```

```
[[3 5 2 4]
```

```
[7 6 8 8]
```

```
[1 6 7 7]]
```

```
3
```

```
1
```

```
7
```

```
[[12 5 2 4]
```

```
[ 7 6 8 8]
```

```
[ 1 6 7 7]]
```

[3 0 3 3 7 9]

INFERENCE:

Array indexing is required for accessing the elements in that array. In the above program I have learnt to implement array indexing using numpy for a three dimensional array.

RESULT: This program was successfully executed using NUMPY.

EX.NO: 3	ARRAY SLICING using NUMPY
DATE:	

AIM:

To write a python program to implement array slicing using numpy

ALGORITHM:

Step1: START

Step2: Import necessary libraries -numpy

Step 3:Using arrange function, print n elements

Step 4:By using the slice method, [x:n] , array slicing can be done

Step 5: Similarly, array slicing for the two dimensional array can be done

Step 6: STOP

PROGRAM:

```
import numpy as np
```

```
np.random.seed(0) # seed for reproducibility
```

```
x1 = np.random.randint(10, size=6) # One-dimensional array
```

```
x2 = np.random.randint(10, size=(3, 4)) # Two-dimensional array
```

```
x3 = np.random.randint(10, size=(3, 4, 5)) # Three-dimensional array
```

```
print(x1[0])
```

```
print(x1[4])
```

```
#To index from the end of the array, negative indices are used
```

```
print(x1[-1])
```

```
print(x1[-2])
```

```
#In a multidimensional array, items are accessed using a comma-separated tuple
```

```
#of indices:
```



```
print(x2) print(x2[0,  
0])  
print(x2[2, 0])  
print(x2[2, -1])
```

```
#modifying values using index notation:
```

```
x2[0, 0] = 12
```

```
print(x2)
```

```
x1[0] = 3.14159 # this will be truncated!
```

```
print(x1)
```

OUTPUT:

```
[5 0 3 3 7 9]
```

```
5
```

```
7
```

```
9
```

```
7
```

```
[[3 5 2 4]
```

```
[7 6 8 8]
```

```
[1 6 7 7]]
```

```
3
```

```
1
```

```
7
```

```
[[12 5 2 4]
```

```
[ 7 6 8 8]
```

```
[ 1 6 7 7]]
```

```
[3 0 3 3 7 9]
```

INFERENCE:

Array slicing is required for accessing certain the elements in that array. In the above program I have learnt to implement array slicing using numpy.

RESULT: This program was successfully executed using NUMPY.

EX.NO:4	SUBARRAYS using NUMPY
DATE:	

AIM:

To write a python program to implement subarrays using numpy

ALGORITHM:

Step 1:START

Step 2:Import the necessary libraries – numpy

Step 3:Usingrandint and random module create a two dimensional arrayStep

4:Extract a n*n subarray from main array

Step 5:print the elements in sub array

Step 6:STOP

PROGRAM:

```
import numpy as np
x = np.arange(10)
print(x)
print(x[:5]) # first five elements
print(x[5:]) # elements after index 5
print(x[4:7]) # middle subarray
print(x[::2]) # every other element
print(x[1::2]) # every other element, starting at index 1
print(x[::-1]) # all elements, reverse
print(x[5::-2]) # reversed every other from index 5
x2 = np.random.randint(10, size=(3, 4)) # Two-dimensional array
print(x2)
print(x2[:2, :3]) # two rows, three columns
print(x2[:3, ::2]) # all rows, every other column
print(x2[::-1, ::-1]) # subarray dimensions reversed together
print(x2[:, 0]) # first column of x2
print(x2[0, :]) # first row of x2
print(x2[0]) # equivalent to x2[0, :]
```

OUTPUT:

```
[0 1 2 3 4 5 6 7 8 9]
```

```
[0 1 2 3 4]
```

```
[5 6 7 8 9]
```

```
[4 5 6]
[0 2 4 6 8]
[1 3 5 7 9]
[9 8 7 6 5 4 3 2 1 0]
[5 3 1]
[[6 4 3 4]
 [8 6 0 6]
 [4 7 2 1]]
[[6 4 3]
 [8 6 0]]
[[6 3]
 [8 0]
 [4 2]]
[[1 2 7 4]
 [6 0 6 8]
 [4 3 4 6]]
[6 8 4]
[6 4 3 4]
[6 4 3 4]
```

INFERENCE:

Sub arrays are required for further processing. From this program, we learnt to extract a subarray from the main two dimensional array and print the elements in sub array

RESULT: This program was successfully executed using NUMPY.

EX.NO:5	DATA INDEXING AND SELECTION USING PANDAS
DATE:	

AIM:

To write a python program to implement data indexing and selection using pandas

ALGORITHM:

- Step 1:START
- Step 2:Import the necessary libraries – pandas
- Step 3:Create a series using series module from pandas
- Step 4:Create a rows and columns (i.e) index and values respectively using pandas series function
- Step 5:print the one dimensional array within a range using string slicing
- Step 6:STOP

PROGRAM:

```
#Subarrays as no-copy views
import numpy as np
x2 = np.random.randint(10, size=(3, 4)) # Two-dimensional array

#extract a 2x2 subarray from this
x2_sub = x2[:2, :2]
print(x2_sub)

#if we modify this subarray, we'll see that the original array is changed!
x2_sub[0, 0] = 99
print(x2_sub)
print(x2)

#when we work with large datasets, we can access and process pieces of these datasets without the
need to copy the underlying data buffer.
```

OUTPUT:

```
[[0 1]
 [8 4]]
[[99 1]
 [ 8 4]]
[[99 1 2 2]]
```

```
[ 8 4 5 9]
```

```
[ 9 3 6 5]]
```

INFERENCE:

Pandas are packages that can be added to python for doing the data analysis. From this program,we learnt to construct series as objects using pandas libraries.

RESULT: This program was successfully executed using PANDAS.

EX.NO:6	OBJECT as Series using PANDAS
DATE:	

AIM:

To write a python program to implement object as series using pandas

ALGORITHM:

- Step 1:START
- Step 2:Import the necessary libraries-numpy,pandas.
- Step 3:Create a series using numpy array
- Step 4:Create a specialized dictionary and build a series.Step
- 5:print the series by using the pandas
- Step 6:STOP

PROGRAM:

```
#PANDAS SERIES AS OBJECT
import numpy as np
import pandas as pd
data = pd.Series([0.25, 0.5, 0.75, 1.0])
print(data)
print(data.values)
print(data.index)
print(data[1])
print(data[1:3])
#series as numpy array
data = pd.Series([0.25, 0.5, 0.75, 1.0],index=['a', 'b', 'c', 'd'])
print(data)
print(data['b'])
data = pd.Series([0.25, 0.5, 0.75, 1.0],index=[2, 5, 3, 7])
print(data)
print(data[5])
#series as specilized dictionary
population_dict = {'California': 38332521,
                  'Texas': 26448193,
                  'New York': 19651127,
                  'Florida': 19552860,
                  'Illinois': 12882135}
population = pd.Series(population_dict)
print(population)
print(population['California'])
```

```

print(population['California':'Florida'])
#constructing series objects
a=pd.Series([2, 4, 6])
print(a)
b=pd.Series(5, index=[100, 200, 300])
print(b)
c=pd.Series({2:'a', 1:'b', 3:'c'})
print(c);
#after indexing
c=pd.Series({2:'a', 1:'b', 3:'c'}, index=[3, 2])
print(c)

```

OUTPUT:

```

0 0.25
1 0.50
2 0.75
3 1.00
dtype: float64
[0.25 0.5 0.75 1.
 ]
RangeIndex(start=0, stop=4,
step=1)0.5
1 0.50
2 0.75
dtype:
float64a
0.25
b 0.50
c 0.75
d 1.00
dtype:
float640.5
2 0.25
5 0.50
3 0.75
7 1.00
dtype:
float640.5
California 38332521
Texas 26448193
New York 19651127
Florida 19552860
Illinois 12882135
dtype:
int64
38332521
California 38332521
Texas 26448193
New York 19651127
Florida 19552860
dtype:
int640
2

```



```
1 4
2 6
dtype: int64
100 5
200 5
300 5
dtype:
int64
2 a
1 b
3 c
dtype: object
3 c
2 a
dtype: object
```

INFERENCE:

Pandas are packages that can be added to python for doing the data analysis. From this problem, we learnt to create a dataframe as specialized dictionary using pandas library functions

RESULT: This program was successfully executed using PANDAS.

EX.NO:7	DATAFRAME OBJECT SERIES AS SPECILIZED DICTIONARY USING PANDAS
DATE:	

AIM:

To write a python program to implement dataframe object series as specilized dictionary usingpandas

ALGORITHM:

Step 1:START
Step 2:Import the necessary libraries-pandas
Step 3:Create a dictionary named population_dict.
Step 4:create a series by using the pandas libraries
Step 5:print the results.
Step 6:STOP

PROGRAM:

```
import pandas as pd
#PANDAS DATAFRAME OBJECT
#series as specilized dictionary
population_dict = {'California': 38332521,
                  'Texas': 26448193,
                  'New York': 19651127,
                  'Florida': 19552860,
                  'Illinois': 12882135}
population = pd.Series(population_dict)
area_dict = {'California': 423967, 'Texas': 695662, 'New York': 141297, 'Florida':
170312, 'Illinois': 149995}
area = pd.Series(area_dict)
print(area)
print()
states = pd.DataFrame({'population': population, 'area': area})
print(states)
print()
print(states.index)
print()
print(states.columns)
print()
#dataframe as specilized dictionary
```

```
print(states['area'])
print()
a= pd.DataFrame(population, columns=['population'])
print(a)
print()
```

OUTPUT:

```
California  423967
Texas       695662
New York    141297
Florida     170312
Illinois    149995
```

```
dtype: int64
```

```
population  area
```

```
California  38332521  423967
Texas       26448193  695662
New York    19651127  141297
Florida     19552860  170312
Illinois    12882135  149995
```

```
Index(['California', 'Texas', 'New York', 'Florida', 'Illinois'], dtype='object')
```

```
Index(['population', 'area'], dtype='object')
```

```
California  423967
Texas       695662
New York    141297
Florida     170312
Illinois    149995
```

```
Name: area, dtype: int64
```

```
population
```

```
California  38332521
Texas       26448193
New York    19651127
Florida     19552860
Illinois    12882135
```

INFERENCE:

Pandas are packages that can be added to python for doing the data analysis. From this problem, we learnt to have to create dataframe object series as specialized dictionary using pandas.

RESULT: This program was successfully executed using PANDAS.

EX.NO:8	KNN CLASSIFICATION FOR USE OF IRIS DATASET
DATE:	

AIM:

To write a python program to implement knn classification for use of iris dataset

ALGORITHM:

Step 1: Load and Train the IRIS data

Step 2: Initialize K to your chosen number of neighbours.

Step 3: For each example in the data

- i. Calculate the distance between the query example and the current example from the data.
- ii. Add the distance and the index of the example to an ordered collection.
- iii. Sort the ordered collection of distances and indices from smallest to largest (in ascending order) by the distances
- iv. Pick the first K entries from the sorted collection
- v. Get the labels of the selected K entries
- vi. Classify the new category as the mode of the K labels and return type

PROGRAM:

```
# Make Predictions with k-nearest neighbors on the Iris Flowers Dataset from csvimport
```

```
reader
from math import sqrt
```

```
# Load a CSV file
def load_csv(filename): dataset = list()
with open(filename, 'r') as file: csv_reader = reader(file) for row in csv_reader:if not
row:
continue dataset.append(row)
return dataset
```

```
# Convert string column to float
def str_column_to_float(dataset, column):for
row in dataset:
row[column] = float(row[column].strip())
```

```
# Convert string column to integer
def str_column_to_int(dataset, column):
class_values = [row[column]
```

```

for row in dataset] unique = set(class_values)
lookup = dict()
for i, value in enumerate(unique): lookup[value] = i print('[%s] =>
%d' % (value, i)) for row in dataset: row[column] =
lookup[row[column]] return lookup
# Find the min and max values for each column def
dataset_minmax(dataset):
minmax = list()
for i in range(len(dataset[0])):
col_values = [row[i] for row in dataset] value_min = min(col_values) value_max =
max(col_values)
minmax.append([value_min, value_max]) return minmax

# Rescale dataset columns to the range 0-1 def
normalize_dataset(dataset, minmax):
for row in dataset:
for i in range(len(row)):
row[i] = (row[i] - minmax[i][0]) / (minmax[i][1] - minmax[i][0])

# Calculate the Euclidean distance between two vectors def
euclidean_distance(row1, row2):
distance = 0.0
for i in range(len(row1)-1):
distance += (row1[i] - row2[i])**2 return sqrt(distance)

# Locate the most similar neighbors
def get_neighbors(train, test_row, num_neighbors): distances = list() for train_row
in train:
dist = euclidean_distance(test_row, train_row)
distances.append((train_row, dist))
distances.sort(key=lambda tup: tup[1]) neighbors = list()
for i in range(num_neighbors): neighbors.append(distances[i][0]) return neighbors

# Make a prediction with neighbors
def predict_classification(train, test_row, num_neighbors): neighbors =
get_neighbors(train, test_row, num_neighbors) output_values = [row[-1] for row in
neighbors]
prediction = max(set(output_values), key=output_values.count) return prediction

# Make a prediction with KNN on Iris Dataset filename = 'iris.csv' dataset =
load_csv(filename) for i in range(len(dataset[0])-1): str_column_to_float(dataset, i)
# convert class column to integers

```

```
str_column_to_int(dataset, len(dataset[0])-1) # define model parameter num_neighbors = 5
# define a new record row = [5.1,3.7,1.5,0.4] # predict
the label
label = predict_classification(dataset, row, num_neighbors)
print('Data=%s, Predicted: %s' % (row, label))
```

OUTPUT:

```
[Setosa] => 0
[Versicolor] => 1
[Virginica] => 2
Data=[5.1, 3.7, 1.5, 0.4],
Predicted: 0
```

INFERENCE:

Classification is used to classify the given data into known groups. In this program we classify the IRIS data.

RESULT: This program was successfully executed.

EX.NO:9

DATE:

CLASSIFICATION USING LINEAR REGRESSION

AIM:

To write a python program to implement classification using linear regression

ALGORITHM:

Step1: Consider a set of values x, y.

Step2: Take the linear set of equation $y = a+bx$.

Step3: Computer value of a, b with respect to the given values, $b = \frac{n\sum xy - (\sum x)(\sum y)}{n\sum x^2 - (\sum x)^2}$, $a = \frac{\sum y - b(\sum x)}{n}$.

Step4: Implement the value of a, b in the equation $y = a+ bx$.

Step5: Regress the value of y for any x.

PROGRAM:

```
import numpy as np
import matplotlib.pyplot as plt
from csv import DictReader
def estimate_coef(x, y):
    # number of observations/points
    n = np.size(x)
    # mean of x and y vector
    m_x, m_y = np.mean(x), np.mean(y)
    # calculating cross-deviation and deviation about x
    SS_xy = np.sum(y*x - n*m_y*m_x)
    SS_xx = np.sum(x*x - n*m_x*m_x) #
    # calculating regression coefficients
    b_1 = SS_xy / SS_xx
    b_0 = m_y - b_1*m_x
    return(b_0, b_1)
def plot_regression_line(x, y, b):
    # plotting the actual points as scatter plot
    plt.scatter(x, y, color = "m",
               marker = "o", s = 30)
    # predicted response vector
    y_pred = b[0] + b[1]*x
    # plotting the regression line
    plt.plot(x, y_pred, color = "g")
```

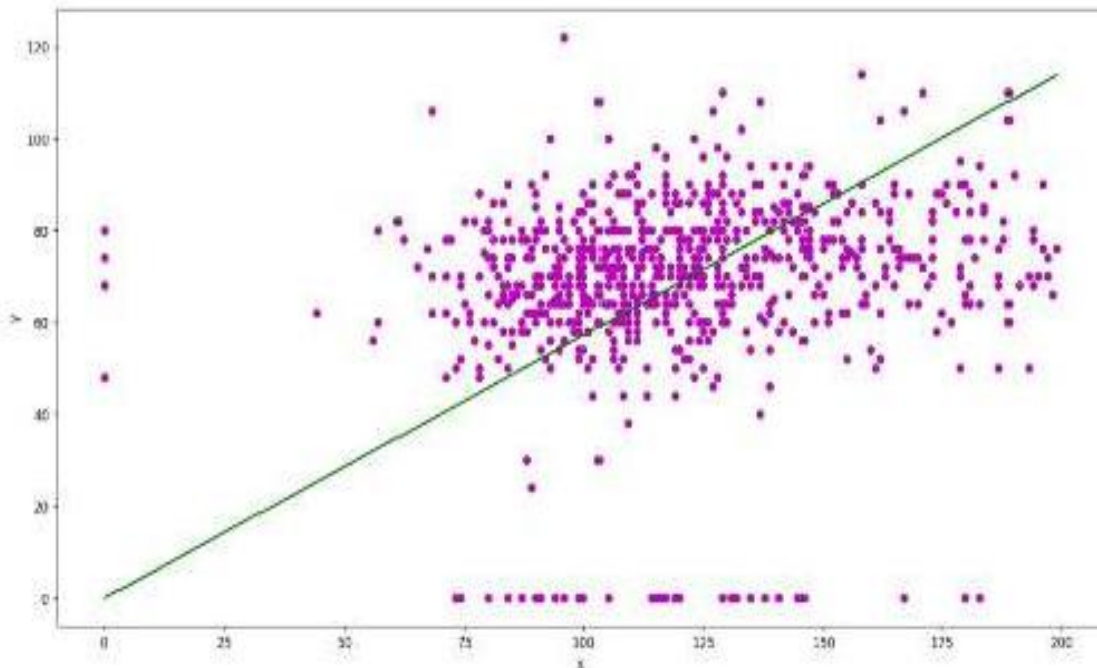


```

# putting labels
plt.xlabel('x')
plt.ylabel('y')
# function to show plot
plt.show()
def main():
    # observations
    Data = []
    X,Y=[],[]
    # opening csv file
    with open('diabetes.csv','r') as file:
        reader = DictReader(file)
        for row in reader:
            Data.append(row)
            for i in Data: X.append(int(i['Glucose']))
            Y.append(int(i['BloodPressure']))
    x = np.array(X)
    y = np.array(Y)
    # estimating coefficients
    b = estimate_coef(x, y)
    print("Estimated coefficients:\nb_0 = {} nb_1 = {}".format(b[0], b[1]))# plotting
    regression_line
    plot_regression_line(x, y, b) if
    name_____=="_main_":
    main()

```

OUTPUT:



INFERENCE:

Linear regression is knowing the relationship between two values .From this program we learnt about the how to implement linear regression using python

RESULT: This program was successfully executed.

EX.NO:10

CLASSIFICATION USING LOGISTIC REGRESSION

DATE:

AIM:

To write a python program to implement classification using logistic regression

ALGORITHM:

Step1: Initialize the variables
Step2: Set the Data frame
Step3: Spilt data set into training and testing. Step4: Fit the data into logistic regression function. Step5: Predict the test data set.
Step6: Print the results.

PROGRAM:

```
import pandas as pd
from sklearn.model_selection import train_test_split
from sklearn.linear_model import LogisticRegression
from sklearn import metrics
import seaborn as sns
import matplotlib.pyplot as plt
from csv import DictReader

Data = []
Glucose, BloodPressure, BMI, Outcome = [], [], [], []
# opening csv file
with open('diabetes.csv', 'r') as file:
    reader = DictReader(file)
    for row in reader:
        Data.append(row)
for i in Data:
    Glucose.append(int(i['Glucose']))
    BloodPressure.append(int(i['BloodPressure']))
    BMI.append(float(i["BMI"]))
    Outcome.append(int(i["Outcome"]))

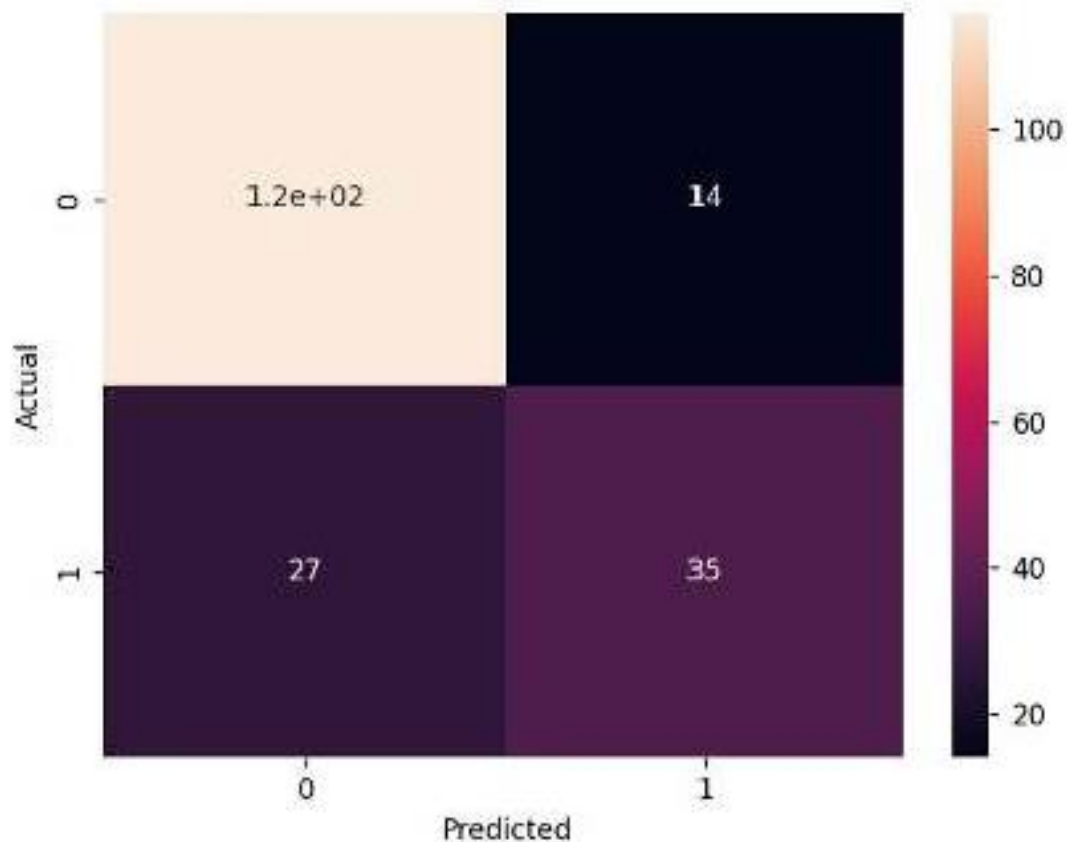
candidates =
{'Glucose': Glucose, 'BMI': BMI, 'BloodPressure': BloodPressure, 'Outcome': Outcome}
```

```

df = pd.DataFrame(candidates,columns= ['Glucose',
'BMI','BloodPressure','Outcome'])
print (df)
print("Df printed\n")
X = df[['Glucose', 'BMI','BloodPressure']]
y = df['Outcome']
X_train,X_test,y_train,y_test =
train_test_split(X,y,test_size=0.25,random_state=0)
print (X_train)
print (y_train)
print("Train\n")
logistic_regression= LogisticRegression()
logistic_regression.fit(X_train,y_train)
y_pred=logistic_regression.predict(X_test)
confusion_matrix = pd.crosstab(y_test, y_pred, rownames=['Actual'],
colnames=['Predicted'])
sn.heatmap(confusion_matrix, annot=True)
print('Accuracy: ',metrics.accuracy_score(y_test, y_pred))
print (X_test) #test dataset
print (y_pred) #predicted values
print('confusion_matrix:', confusion_matrix, sep='\n', end='\n\n')
plt.show()

```

OUTPUT:



INFERENCE:

Logistic regression is an example of supervised learning. It is used to calculate or predict the probability of a binary (yes/no) event occurring. From this program we learnt to draw the logistics regressions using python

RESULT: This program was successfully executed.

EX.NO:12	MULTIPLE REGRESSION ANALYSIS
DATE:	

AIM:

To write a python program to implement multiple regression analysis

ALGORITHM:

Step1: Get the multi-attribute dataset using the Scikit-learn data source.

Step 2: Create a regression object.

Step 3: Train the dataset with the regression model fit.

Step 4: Get and print the regression coefficients and variance.

Step 5. Plot the residual error.

PROGRAM:

```
import matplotlib.pyplot as plt
import numpy as np
from sklearn import datasets, linear_model, metrics # load the
boston dataset

boston = datasets.load_boston(return_X_y=False) # defining
feature matrix(X) and response vector(y)

X = boston.data
y = boston.target

# splitting X and y into training and testing sets from
sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.4,
random_state=1)

# create linear regression object

reg = linear_model.LinearRegression()

# train the model using the training sets
reg.fit(X_train, y_train)

# regression coefficients
print('Coefficients: ', reg.coef_)
```

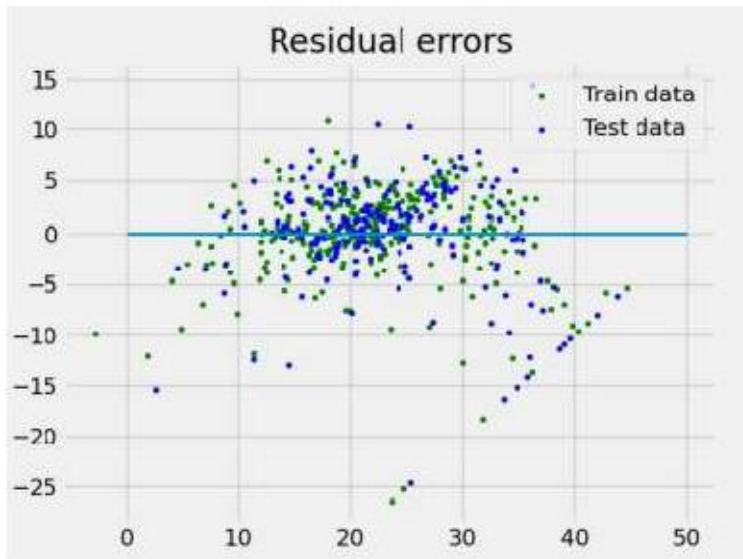
```
# variance score: 1 means perfect prediction print('\nVariance score:
{}'.format(reg.score(X_test, y_test))) # plot
for residual error
## setting plot style plt.style.use('fivethirtyeight')
## plotting residual errors in training data plt.scatter(reg.predict(X_train),reg.predict(X_train) - y_train,
color = "green", s = 10, label = 'Train data')
## plotting residual errors in test data plt.scatter(reg.predict(X_test),reg.predict(X_test) - y_test,
color = "blue", s = 10, label = 'Test data')##
plotting line for zero residual error
plt.hlines(y = 0, xmin = 0, xmax = 50, linewidth = 2)##
plotting legend plt.legend(loc = 'upper right') ## plot title
plt.title("Residual errors")
## method call for showing the plot plt.show()
```

OUTPUT:

Coefficients:

```
[-8.95714048e-02 6.73132853e-02
 5.04649248e-02 2.18579583e+00
 -1.72053975e+01 3.63606995e+00
 2.05579939e-03 -1.36602886e+00
 2.89576718e-01 -1.22700072e-02 -
 8.34881849e-01 9.40360790e-03
 -5.04008320e-01]
```

Variance score: 0.720905667266178



INFERENCE:

Multiple regression is a statistical technique that can be used to analyze the relationship between a single dependent variable and several independent variables. The objective of multiple regression analysis is to use the independent variables whose values are known to predict the value of the single dependent value. From this program we learnt to draw the multiple linear regression.

RESULT: This program was successfully executed.

EX.NO: 13	NORMAL CURVES
DATE:	

AIM:

To write a python program to implement normal curves

ALGORITHM:

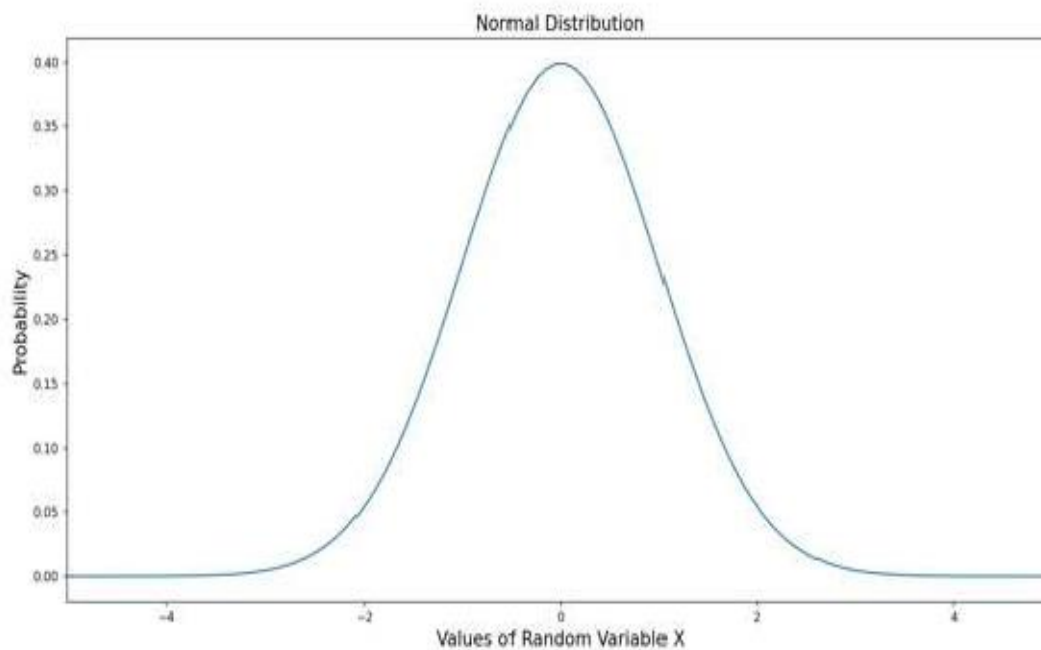
- Step1: Set the Mean as 0 and Standard Deviation as 1.
- Step2: Generate the set x of 100 random numbers in the range of -5 to 5.
- Step3: Define the probability density function using x.
- Step4: Plot the Normal Distribution.

PROGRAM:

```
import numpy as np
import matplotlib.pyplot as plt
from scipy import stats

# Create a standard normal distribution with mean as 0 and standard deviation as
1
#
mu = 0
std = 1
snd = stats.norm(mu, std)
#
# Generate 100 random values between -5, 5
#
x = np.linspace(-5, 5, 100)
#
# Plot the standard normal distribution for different values of random variable
# falling in the range -5, 5
#
plt.figure(figsize=(7.5,7.5))
plt.plot(x, snd.pdf(x))
plt.xlim(-5, 5)
plt.title('Normal Distribution', fontsize='15')
plt.xlabel('Values of Random Variable X', fontsize='15')
plt.ylabel('Probability', fontsize='15')
plt.show()
```

OUTPUT:



INFERENCE:

Normal distribution, also known as the Gaussian distribution, is a probability distribution that is symmetric about the mean, showing that data near the mean are more frequent in occurrence than data far from the mean. In graphical form, the normal distribution appears as a "bell curve". From this program we learnt to draw a curve for normal distribution using matplotlib and numpy functions

RESULT: This program was successfully executed.

EX.NO: 14

CORRELATION ANALYSIS

DATE:

AIM:

To write a python program to implement correlation analysis

ALGORITHM:

Step1: Compute the value of \bar{x} & \bar{y} .

Step 2: Compute $\sum_{i=1}^n (X - \bar{x})(Y - \bar{y})$

Step 3: Compute

$$r_{x,y} = \frac{\sum_{i=1}^n (x_i - \bar{x})(y_i - \bar{y})}{\sqrt{\sum_{i=1}^n (x_i - \bar{x})^2} \sqrt{\sum_{i=1}^n (y_i - \bar{y})^2}}$$

Step 4: Find it is highly correlated or low correlated and display the result.

PROGRAM:

```
# Python Program to find correlation coefficient. import math
# function that returns correlation coefficient. def correlationCoefficient(X, Y, n ):sum_X = 0
    sum_Y = 0
    sum_XY = 0
    squareSum_X = 0
    squareSum_Y = 0
    i = 0
    while i<n :
        # sum of elements of array X. sum_X = sum_X + X[i]
```

```

# sum of elements of array Y. sum_Y = sum_Y + Y[i]

# sum of X[i] * Y[i].
sum_XY = sum_XY + X[i] * Y[i]

# sum of square of array elements. squareSum_X = squareSum_X + X[i] *X[i]
squareSum_Y = squareSum_Y + Y[i] * Y[i]

i = i + 1

# use formula for calculating correlation # coefficient.
corr = (float)(n * sum_XY - sum_X * sum_Y)/ ( float )(math.sqrt((n *squareSum_X
-sum_X * sum_X)* (n * squareSum_Y - sum_Y * sum_Y))) return corr

# Driver function
X = [15, 18, 21, 24, 27]
Y = [25, 25, 27, 31, 32]

print(X) print(Y)

# Find the size of array. n = len(X)

# Function call to correlationCoefficient. z = correlationCoefficient(X, Y, n)if(abs(z) > 0.5):
print ('{0:.6f}'.format(z), "Highly COrelated") else:print('{0:.6f}'.format(z),"Low
Correlated")

```

OUTPUT:

[15, 18, 21, 24, 27]

[25, 25, 27, 31, 32]

0.953463 Highly Correlated

INFERENCE:

Correlation is a statistical measure that expresses the extent to which two variables are linearly related (meaning they change together at a constant rate). It's a common tool for describing simple relationships without making a statement about cause and effect. From this program we learnt about the correlation analysis technique using python

RESULT: This program was successfully executed.

EX.NO: 15	MEAN,MEDIAN, MODE, STANDARD DEVIATION
DATE:	

AIM:

To write a python program to implement mean, median, mode and standard deviation.

ALGORITHM:

Step1: Take a list of 8 Numbers.

Step2: Compute the Mean value by simple Computation and print it.

Step3: Compute the Mean value using numpy method and print it.

Step4: Compute the Median value by simple Computation and print it.

Step5: Compute the Mode value by simple Computation and print it.

Step6: Compute the Mode value using numpy method and print it.

Step7: Compute the Standard Deviation by simple Computation and print it.

Step8: Compute the Standard Deviation using Numpy and print it.

PROGRAM:

```
# Write a program to compute mean, median, mode and Standard Deviation
import numpy as np
from collections import Counter
from scipy import stats
# Finding Mean by simple Computation
a=[11, 21, 34, 22, 27, 11, 23, 21]
mean = sum(a)/len(a)
print("Finding Mean by simple Computation")
print(mean)
# Finding Mean using numpy method
mean = np.mean(a)
print("Finding Mean using numpy method ")
print(mean)
#Finding Median by simple Computation.
def median(nums):
    nums.sort()
    if len(nums)%2 == 0:
        return int((nums[len(nums)//2-1]+nums[len(nums)//2])/2)
    else:
        return nums[len(nums)//2]
print("Finding Median by simple Computation")
print(median(a))
print("Finding Median by numpy method")
```

```
print(np.median(a))
# Finding Mode by simple Computation
data = dict(Counter(a))
mode = [k for k, v in data.items() if v == max(list(data.values()))]
print("Finding Mode by simple Computation ")
print (mode)
# Finding Mode using numpy method
print("Finding Mode using numpy method")
print (stats.mode(a,axis=None,keepdims=True))
# Find Standard deviation by simple computation
n=len(a)
std=(sum(map(lambda x: (x-sum(a)/n)**2,a)/n)**0.5)
print(std)
# Find Standard deviation using numpy method
print(np.std(a))
```

OUTPUT:

```
Finding Mean by simple Computation
21.25
Finding Mean using numpy method
21.25
Finding Median by simple Computation21.5
Finding Median by numpy method
21.5
Finding Mode by simple Computation[11,
21]
Finding Mode using numpy method
ModeResult(mode=array([11]), count=array([2]))
7.1545440106270926
7.1545440106270926
```

INFERENCE:

Mean, median, mode and standard deviation are used for data analysis in data science. From this program we have learnt how to calculate Mean, median, mode and standard deviation using simple method and numpy method.

RESULT: This program was successfully executed

EX.NO: 14	DATA VISUALIZATION
DATE:	

AIM:

To write a python program to implement data visualization

ALGORITHM:

- Step1: Load the IRIS Dataset and Wine Review Dataset
- Step 2: Create the Color Scatter Plot of IRIS Dataset.
- Step 3: Create the Line chart for each attributes of IRIS Dataset.
- Step 4: Create the Histogram for Wine Review Scores.
- Step 5: Create the Bar Chart for Wine Review Scores.
- Step 6: Create the multiple histogram for attributes of IRIS Dataset.
- Step 7: Create the vertical bar chart for Wine Review Scores using plot.bar().
- Step 8: Create the horizontal bar chart for Wine Review Scores using plot.bar().
- Step 9: Create the bar chart for Wine Review with highest cost five different Counties.

PROGRAM:

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

iris = pd.read_csv('iris.csv', names=['sepal_length', 'sepal_width', 'petal_length', 'petal_width', 'class'])
print(iris.head())

wine_reviews = pd.read_csv('winemag-data-130k-v2.csv', index_col=0)
wine_reviews.head()

# Create Color Scatter Plotting
colors = {'Iris-setosa': 'r', 'Iris-versicolor': 'g', 'Iris-virginica': 'b'} # create a figure and axis
fig, ax = plt.subplots() # plot each data-point
for i in range(len(iris['sepal_length'])):
```



```
ax.scatter(iris['sepal_length'][i], iris['sepal_width'][i],color=colors[iris['class'][i]])# set a title and labels
ax.set_title('Iris Dataset') ax.set_xlabel('sepal_length')
ax.set_ylabel('sepal_width')plt.show()
```

```
# Create Line Chart Plotting columns = iris.columns.drop(['class']) # create x data
ax_data = range(0, iris.shape[0]) # create figure and axis
fig, ax = plt.subplots() # plot each column
for column in columns:
    ax.plot(ax_data, iris[column], label=column) # set title and legend
ax.set_title('Iris Dataset') ax.legend()
plt.show()
```

```
# create figure and axis fig, ax = plt.subplots() # plot histogram
ax.hist(wine_reviews['points']) # set title and labels
ax.set_title('Wine Review Scores') ax.set_xlabel('Points')
ax.set_ylabel('Frequency') plt.show()
```

```
# create a figure and axis fig, ax = plt.subplots()# count the occurrence of each class
data = wine_reviews['points'].value_counts() # get x and y data
points = data.index frequency = data.values # create bar chart ax.bar(points,frequency) # set title and labels
ax.set_title('Wine Review Scores') ax.set_xlabel('Points')
ax.set_ylabel('Frequency') plt.show()
```

```
iris.plot.hist(subplots=True, layout=(2,2), figsize=(10, 10), bins=20) plt.show()
```

```
wine_reviews['points'].value_counts().sort_index().plot.bar() plt.show()
```

```
wine_reviews['points'].value_counts().sort_index().plot.barh() plt.show()
```

```
wine_reviews.groupby("country").price.mean().sort_values(ascending=False)[:5  
].plot.bar()  
plt.show()
```

```
# Correlation Matrix corr = iris.corr() fig, ax =  
plt.subplots() # create heatmap  
ax.imshow(corr.values)
```

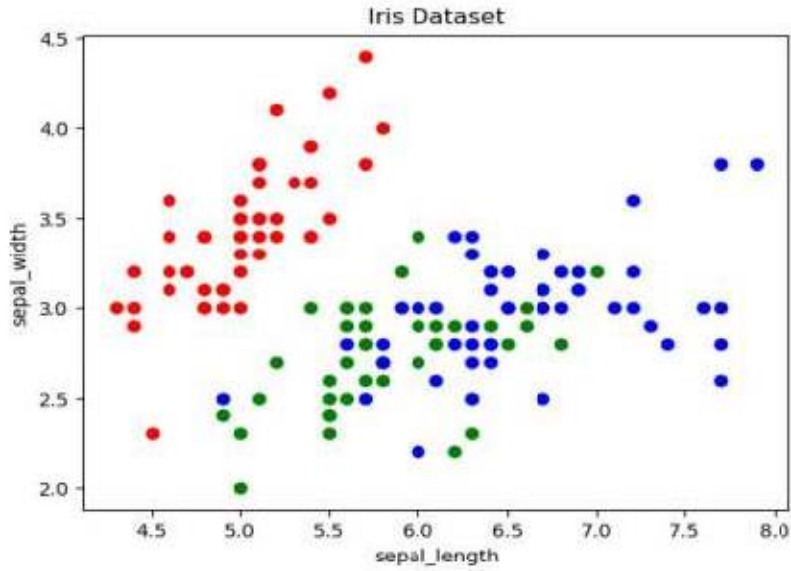
```
# set labels ax.set_xticks(np.arange(len(corr.columns)))  
ax.set_yticks(np.arange(len(corr.columns))) ax.set_xticklabels(corr.columns)  
ax.set_yticklabels(corr.columns)
```

```
# Rotate the tick labels and set their alignment. plt.setp(ax.get_xticklabels(),rotation=45,  
ha="right",  
rotation_mode="anchor")
```

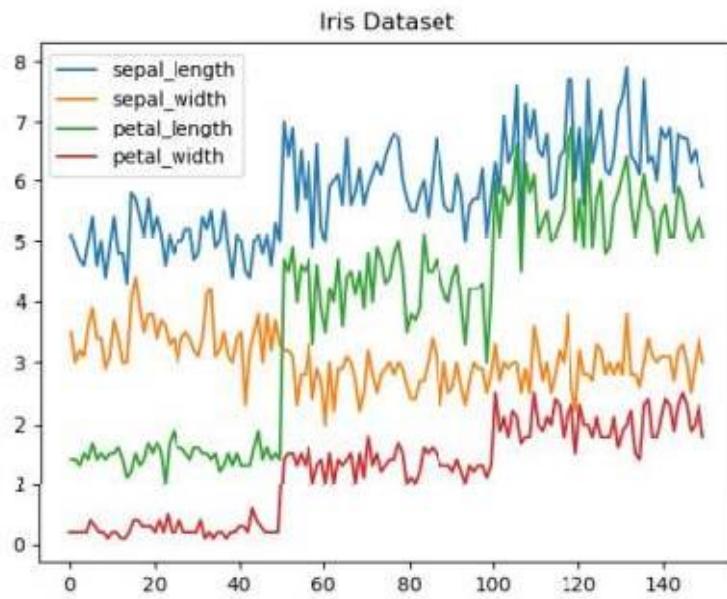
```
# Loop over data dimensions and create text annotations. for i in  
range(len(corr.columns)):  
for j in range(len(corr.columns)):  
text = ax.text(j, i, np.around(corr.iloc[i, j], decimals=2), ha="center",  
va="center", color="black")  
plt.show()
```

OUTPUT:

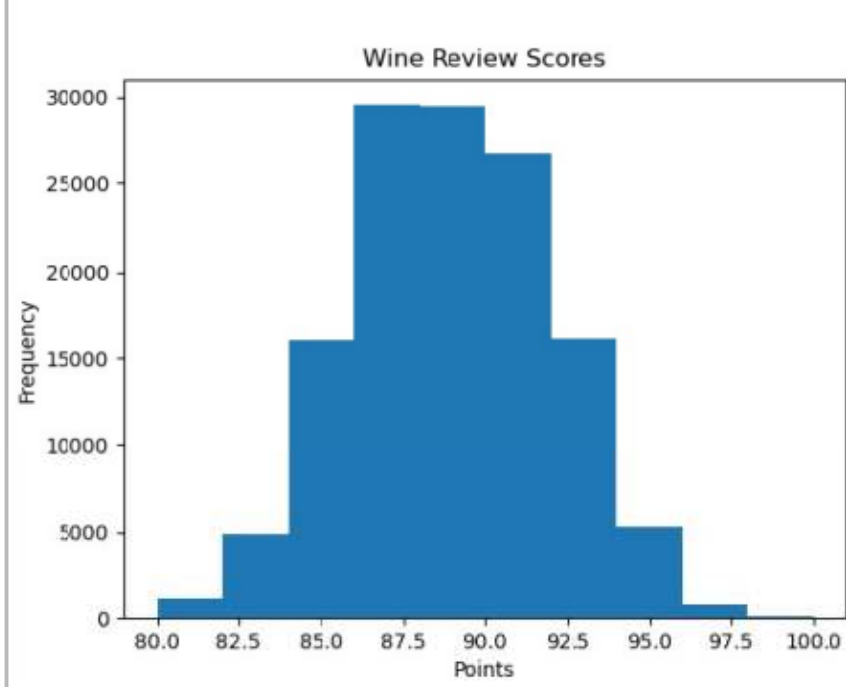
Scatter Plot of IRIS Dataset



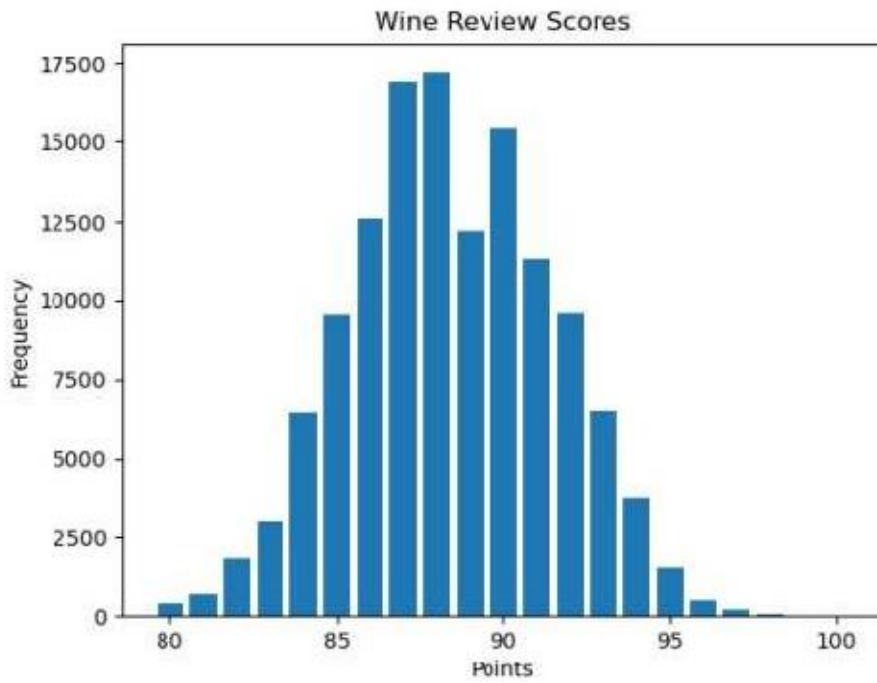
Line chart for each attribute of IRIS Dataset



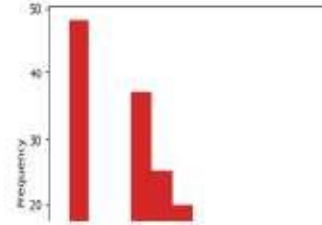
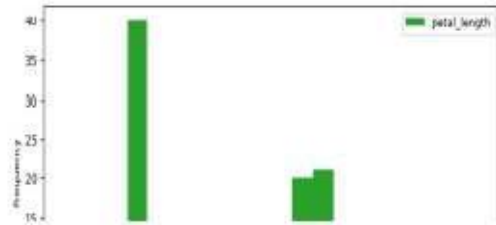
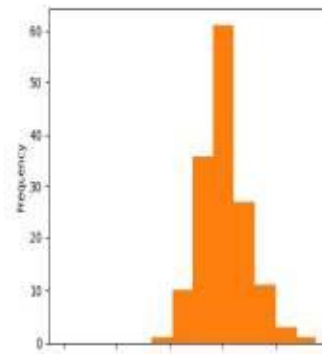
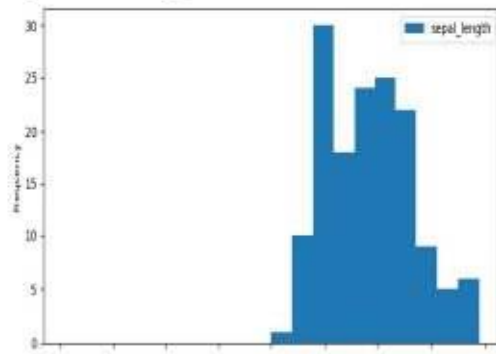
Histogram for Wine Review Scores.



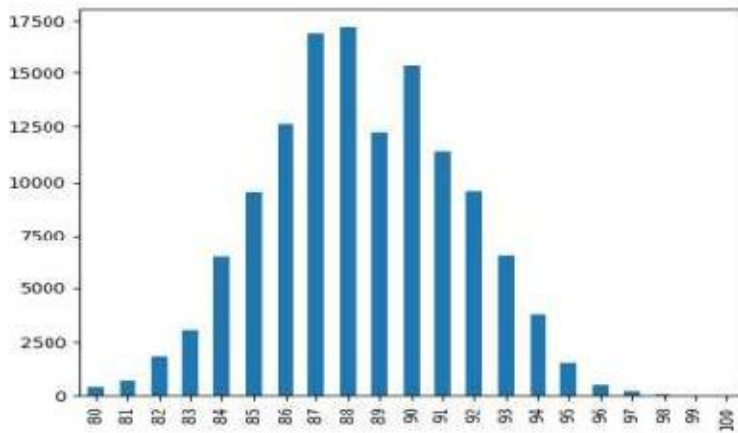
Bar Chart for Wine Review Scores



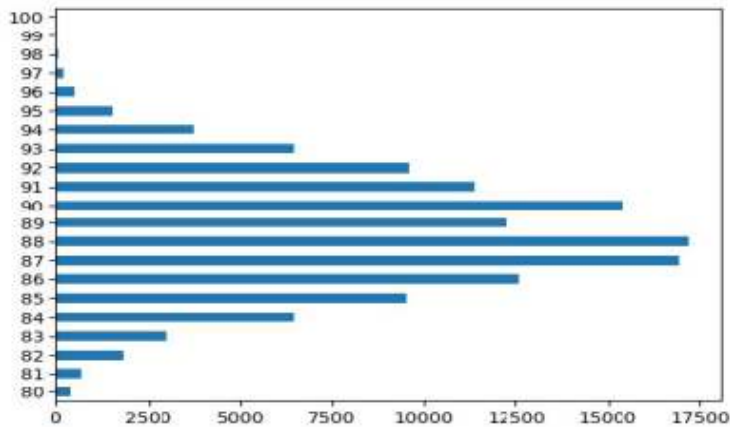
Multiple histogram for attributes of IRIS Dataset



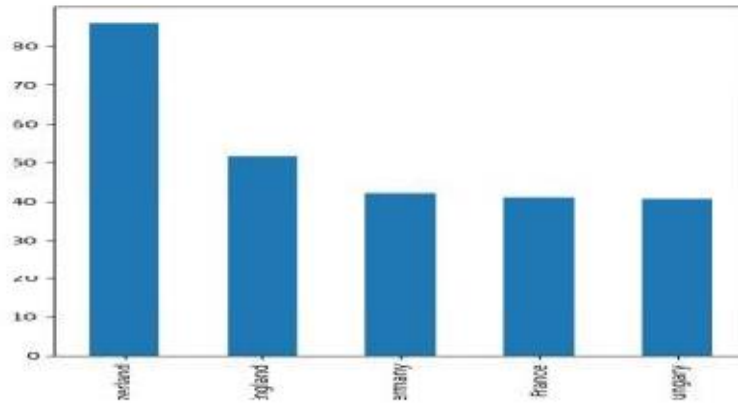
Vertical bar chart for Wine Review Scores



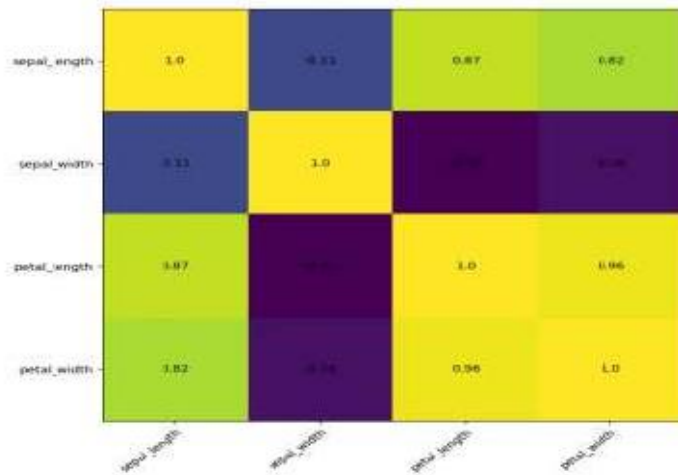
Horizontal bar chart for Wine Review Score



Bar chart for Wine Review with highest cost five different Counties.



Correlation Matrix



INFERENCE:

Data visualization is a way to represent information graphically, highlighting patterns and trends in data and helping the reader to achieve quick insights. From this program we learnt how to visualize data using python.

RESULT: This program was successfully executed.

CONTENT BEYOND SYLLABUS

EX.NO: 15	PRINCIPAL COMPONENT ANALYSIS
DATE:	

AIM:

To write a python Application Program to demonstrate the Principal Component Analysis.

ALGORITHM:

Step 1: Get data.

Step 2: Compute the mean vector (μ). Step 3:

Subtract mean from the given data. Step 4:

Calculate the covariance matrix.

Step 5: Calculate the eigen vectors and eigen values of the covariance matrix. Step

6: Choosing components and forming a feature vector.

Step 7: Deriving the new data set.

PROGRAM:

```
import matplotlib.pyplot as plt

import pandas as pd

import numpy as np

import seaborn as sns

from sklearn.datasets import load_breast_cancer

cancer = load_breast_cancer()

cancer.keys()

df = pd.DataFrame(cancer['data'], columns=cancer['feature_names'])

df.head()

from sklearn.preprocessing import StandardScaler

scaler = StandardScaler()
```

```

scaler.fit(df)

scaled_data = scaler.transform(df)

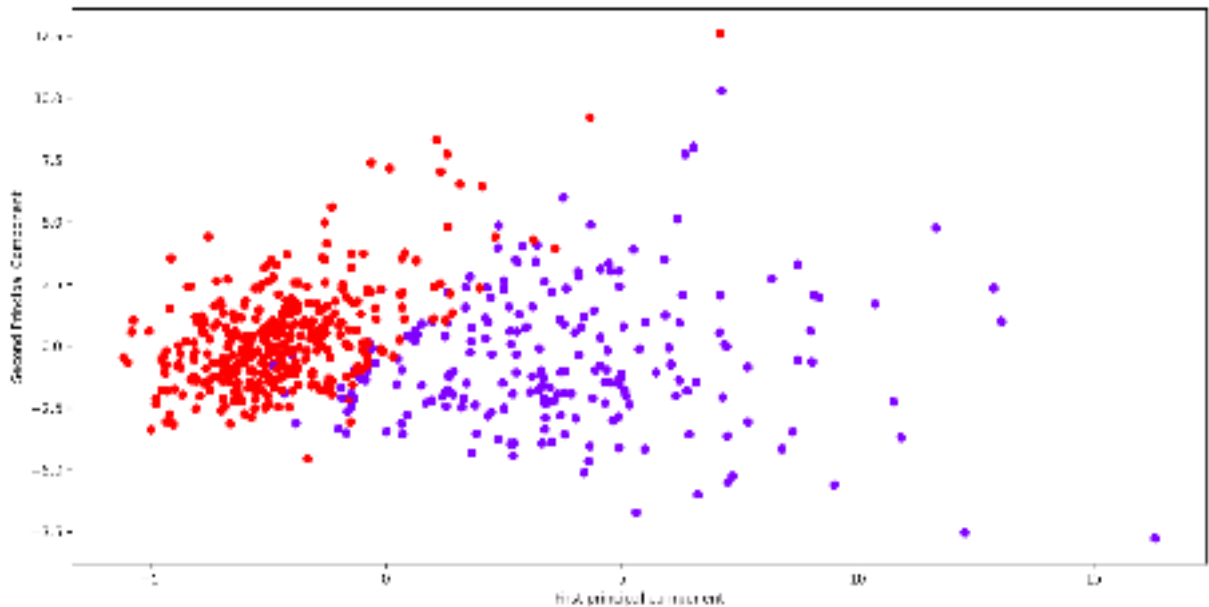
from sklearn.decomposition import PCA

pca = PCA(n_components=2)

pca.fit(scaled_data)
x_pca = pca.transform(scaled_data)
print("Actual size",scaled_data.shape)
print("After PCA",x_pca.shape)
plt.figure(figsize=(8,6))
plt.scatter(x_pca[:,0],x_pca[:,1],c=cancer['target'],cmap='rainbow') plt.xlabel('First
principal component')
plt.ylabel('Second Principal Component')
plt.show()
map= pd.DataFrame(pca.components_,columns=cancer['feature_names'])
plt.figure(figsize=(12,6))
sns.heatmap(map,cmap='twilight')plt.show()

```

OUTPUT:



INFERENCE:

Principal components analysis (PCA) is a dimensionality reduction technique that enables you to identify correlations and patterns in a data set so that it can be transformed into a data set of significantly lower dimension without loss of any important information. From this program we learnt how to implement a PCA using python.

RESULT: This program was successfully executed.