```
In [5]: import pandas as pd
 In [9]: | df = pd.read csv('iris.data')
         df.head()
 Out[91:
                            species
             wa la
                    sl sw
          0 5.1 3.5 1.4 0.2 Iris-setosa
          1 4.9 3.0 1.4 0.2 Iris-setosa
          2 4.7 3.2 1.3 0.2 Iris-setosa
          3 4.6 3.1 1.5 0.2 Iris-setosa
          4 5.0 3.6 1.4 0.2 Iris-setosa
In [10]: df.isnull().sum()
Out[10]: pl
                     0
         pw
                     0
                     0
         sl
                     0
         species
                     0
         dtype: int64
In [11]: x = df.drop(columns='species',axis=1)
In [13]: y = df['species']
In [10]: from sklearn.model selection import train test split
         from sklearn.linear_model import LogisticRegression
In [44]: x train , x test, y train, y test = train test split(x,y,test size=0.3, random sta
In [45]: model = LogisticRegression()
In [46]: model.fit(x train,y train)
         pred y = model.predict(x test)
         print(pred_y)
         ['Iris-setosa' 'Iris-versicolor' 'Iris-versicolor' 'Iris-setosa'
           'Iris-virginica' 'Iris-versicolor' 'Iris-virginica' 'Iris-setosa'
           'Iris-setosa' 'Iris-virginica' 'Iris-versicolor' 'Iris-setosa'
           'Iris-virginica' 'Iris-versicolor' 'Iris-versicolor' 'Iris-setosa'
           'Iris-versicolor' 'Iris-versicolor' 'Iris-setosa' 'Iris-setosa'
           'Iris-versicolor' 'Iris-versicolor' 'Iris-virginica' 'Iris-setosa'
           'Iris-virginica' 'Iris-versicolor' 'Iris-setosa' 'Iris-setosa'
           'Iris-versicolor' 'Iris-virginica' 'Iris-versicolor' 'Iris-virginica'
           'Iris-versicolor' 'Iris-virginica' 'Iris-virginica' 'Iris-setosa'
           'Iris-versicolor' 'Iris-setosa' 'Iris-versicolor' 'Iris-virginica'
           'Iris-virginica' 'Iris-setosa' 'Iris-virginica' 'Iris-virginica'
           'Iris-versicolor']
         /home/student/anaconda3/lib/python3.9/site-packages/sklearn/linear model/ logisti
         c.py:814: ConvergenceWarning: lbfqs failed to converge (status=1):
         STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.
         Increase the number of iterations (max_iter) or scale the data as shown in:
             https://scikit-learn.org/stable/modules/preprocessing.html (https://scikit-le
         arn.org/stable/modules/preprocessing.html)
         Please also refer to the documentation for alternative solver options:
              https://scikit-learn.org/stable/modules/linear_model.html#logistic-regression
         (https://scikit-learn.org/stable/modules/linear model.html#logistic-regression)
           n_iter_i = _check_optimize_result(
```

```
In [2]: from sklearn.metrics import accuracy score
In [48]: | acc = accuracy_score(y_test,pred_y)
         print(acc)
         0.977777777777777
In [49]: model.score(x_test,y_test)
Out[49]: 0.977777777777777
In [50]: from sklearn.preprocessing import StandardScaler
In [51]: #std = StandardScaler()
         #x test = std.fit transform(x test)
In [52]: # y test = std.fit transform(y test)
In [53]: # gauzy(input = noramlly distributed) ,
         # bernaulli(y/n.1/0) ,naivy(continuus) , multinomial naivy(categorical),
In [13]: from sklearn.naive bayes import GaussianNB
In [55]: model1 = GaussianNB()
In [56]: NBmodel = model1.fit(x_train,y_train)
         y pred1 = model1.predict(x test)
In [57]: | acc1 = accuracy_score(y_test,y_pred1)
In [58]: print(acc1)
         0.9333333333333333
In [84]: model1.predict([[5.1,3.5,1.4,0.2]])
         /home/student/anaconda3/lib/python3.9/site-packages/sklearn/base.py:450: UserWarn
         ing: X does not have valid feature names, but GaussianNB was fitted with feature
         names
           warnings.warn(
Out[84]: array(['Iris-setosa'], dtype='<U15')</pre>
In [63]: from sklearn.metrics import confusion_matrix
         from sklearn.metrics import ConfusionMatrixDisplay
In [20]: | from sklearn.metrics import recall_score ,precision_score
In [72]: |print("recall Score : ", recall_score(y_test,pred_y,average='micro'))
         In [69]: from sklearn.metrics import classification report
```

```
In [70]: print(classification report(pred y,y test))
                           precision
                                        recall f1-score
                                                            support
             Iris-setosa
                                1.00
                                          1.00
                                                     1.00
                                                                 14
         Iris-versicolor
                                0.94
                                          1.00
                                                     0.97
                                                                 17
          Iris-virginica
                                1.00
                                          0.93
                                                     0.96
                                                                 14
                                                     0.98
                                                                 45
                accuracy
                                0.98
                                          0.98
                                                                 45
               macro avg
                                                     0.98
                                                                 45
                                0.98
                                          0.98
                                                     0.98
            weighted avg
In [77]: new data = [[5.5,3.5,1.2,0.2],[6.7,3.4,4.3,1.2],[6.2,2.6,5.8,1.8]]
         new peredict = NBmodel.predict(new data)
         print(new_peredict)
         ['Iris-setosa' 'Iris-versicolor' 'Iris-virginica']
         /home/student/anaconda3/lib/python3.9/site-packages/sklearn/base.py:450: UserWarn
         ing: X does not have valid feature names, but GaussianNB was fitted with feature
         names
           warnings.warn(
In [81]:
        new_data = [[6.7, 3.4, 4.3, 1.2]]
         new peredict = NBmodel.predict(new data)
         print(new_peredict)
         ['Iris-versicolor']
         /home/student/anaconda3/lib/python3.9/site-packages/sklearn/base.py:450: UserWarn
         ing: X does not have valid feature names, but GaussianNB was fitted with feature
           warnings.warn(
In [831:
         NBmodel.predict([[5.1,3.5,1.4,0.2]])
         /home/student/anaconda3/lib/python3.9/site-packages/sklearn/base.py:450: UserWarn
         ing: X does not have valid feature names, but GaussianNB was fitted with feature
         names
           warnings.warn(
```

Out[83]: array(['Iris-setosa'], dtype='<U15')</pre>

naivy based alogrithm for diabetes dataset

```
In [6]: | df = pd.read csv('diabetes.csv')
          df.head()
Out[6]:
                          Glucose BloodPressure SkinThickness Insulin BMI DiabetesPedigreeFunction Age Outcom
              Pregnancies
           0
                       6
                              148
                                              72
                                                            35
                                                                    0 33.6
                                                                                               0.627
                                                                                                      50
                       1
                                                            29
                                                                    0 26.6
                                                                                               0.351
           1
                               85
                                              66
                                                                                                      31
           2
                       8
                                                             0
                                                                    0 23.3
                                                                                                      32
                              183
                                              64
                                                                                               0.672
                                                            23
                                                                   94 28.1
           3
                       1
                               89
                                              66
                                                                                               0.167
                                                                                                      21
                       0
                              137
                                              40
                                                                   168 43.1
                                                                                               2.288
                                                                                                      33
In [7]: | x = df.drop('Outcome',axis=1)
```

```
In [8]: y = df['Outcome']
In [11]: x_train , x_test, y_train, y_test = train_test_split(x,y,test_size=0.3,random_stat)
In [14]: model2 = GaussianNB()
In [15]: dia_NBmodel = model2.fit(x_train,y_train)
In [16]: pred y3 = model2.predict(x test)
In [17]: predict_this_data = [[6,148,72,35,0,33.6,0.627,50],[2,85,63,22,0,26.6,0.351,29]]
         predicted = dia NBmodel.predict(predict this data)
         print(predicted)
         [10]
         /home/student/anaconda3/lib/python3.9/site-packages/sklearn/base.py:450: UserWarn
         ing: X does not have valid feature names, but GaussianNB was fitted with feature
         names
           warnings.warn(
In [19]: dia_acc = accuracy_score(y_test,pred_y3)
         print(dia acc)
         0.7835497835497836
In [21]: print("Diabetes Recall Score : ",recall_score(pred_y3,y_test,average='micro'))
         Diabetes Recall Score : 0.7835497835497836
In [ ]:
In [ ]:
```