WEEK - 08 - MLP

1. Use Pima Indians Diabetes dataset.

Implement MLP. Use the sigmoidal activation function used in the algorithm in your text book. Fix the number of hidden neurons based on experimentation. With the same number of hidden neurons experiment using three different activation functions.

```
Definition of class MLP:-
import numpy as np
def tanh(z):
   return (np.exp(z) - np.exp(-z)) / (np.exp(z) + np.exp(-z))
class mlp:
   """A Multi-Layer Perceptron"""
   def __init__(
       self, inputs, targets, nhidden, beta=1, momentum=0.9, outtype="logistic"
   ):
       """Constructor"""
       self.nin = np.shape(inputs)[1]
       self.nout = np.shape(targets)[1]
       self.ndata = np.shape(inputs)[0]
       self.nhidden = nhidden
       self.beta = beta
       self.momentum = momentum
       self.outtype = outtype
       # Initialise network
       self.weights1 = (
           (np.random.rand(self.nin + 1, self.nhidden) - 0.5) * 2 / np.sqrt(self.nin)
       self.weights2 = (
           (np.random.rand(self.nhidden + 1, self.nout) - 0.5)
           / np.sqrt(self.nhidden)
   def earlystopping(
       self, inputs, targets, valid, validtargets, eta, niterations=10000
   ):
       valid = np.concatenate((valid, -np.ones((np.shape(valid)[0], 1))), axis=1)
       old_val_error1 = 100002
       old_val_error2 = 100001
       new_val_error = 100000
       print("No. of neurons in hidden layers = ", self.nhidden)
       while ((old_val_error1 - new_val_error) > 0.001) or (
           (old_val_error2 - old_val_error1) > 0.001
           self.mlptrain(inputs, targets, eta, niterations)
           old_val_error2 = old_val_error1
           old_val_error1 = new_val_error
           validout = self.mlpfwd(valid)
           new_val_error = 0.5 * np.sum((validtargets - validout) ** 2)
```

```
print("Stopped, error = ", new val error)
   return new_val_error
def mlptrain(self, inputs, targets, eta, niterations):
   """Train the neural network"""
   # Add the inputs that match the bias node
   inputs = np.concatenate((inputs, -np.ones((self.ndata, 1))), axis=1)
   change = range(self.ndata)
   updatew1 = np.zeros((np.shape(self.weights1)))
   updatew2 = np.zeros((np.shape(self.weights2)))
   for n in range(niterations):
       self.outputs = self.mlpfwd(inputs)
       error = 0.5 * np.sum((self.outputs - targets) ** 2)
       # Different types of output neurons and their activation functions
       if self.outtype == "linear":
           deltao = (self.outputs - targets) / self.ndata
       elif self.outtype == "logistic":
           deltao = (
              self.beta
              * (self.outputs - targets)
              * self.outputs
              * (1.0 - self.outputs)
           )
       elif self.outtype == "softmax":
           deltao = (
              (self.outputs - targets)
              * (self.outputs * (-self.outputs) + self.outputs)
              / self.ndata
           )
       elif self.outtype == "tanh":
           deltao = (
              (self.outputs - targets)
              * (1.0 - np.power(self.outputs, 2))
       else:
          print("error")
       # hidden network delta
       deltah = (
          self.hidden
           * self.beta
           * (1.0 - self.hidden)
           * (np.dot(deltao, np.transpose(self.weights2)))
       )
       updatew1 = (
           eta * (np.dot(np.transpose(inputs), deltah[:, :-1]))
           + self.momentum * updatew1
       updatew2 = (
          eta * (np.dot(np.transpose(self.hidden), deltao))
           + self.momentum * updatew2
       self.weights1 -= updatew1
       self.weights2 -= updatew2
       return error
```

```
def mlpfwd(self, inputs):
   """Run the network forward"""
   self.hidden = np.dot(inputs, self.weights1)
   self.hidden = 1.0 / (1.0 + np.exp(-self.beta * self.hidden))
   self.hidden = np.concatenate(
       (self.hidden, -np.ones((np.shape(inputs)[0], 1))), axis=1
   outputs = np.dot(self.hidden, self.weights2)
   # Different types of output neurons
   if self.outtype == "linear":
       return outputs
   elif self.outtype == "logistic":
       return 1.0 / (1.0 + np.exp(-self.beta * outputs))
   elif self.outtype == "softmax":
       normalisers = np.sum(np.exp(outputs), axis=1) * np.ones(
           (1, np.shape(outputs)[0])
       return np.transpose(np.transpose(np.exp(outputs)) / normalisers)
   elif self.outtype == "tanh":
       return tanh(outputs)
   else:
       print("error")
def confmat(self, inputs, targets):
   """Confusion matrix""
   # Add the inputs that match the bias node
   inputs = np.concatenate((inputs, -np.ones((np.shape(inputs)[0], 1))), axis=1)
   outputs = self.mlpfwd(inputs)
   nclasses = np.shape(targets)[1]
   if nclasses == 1:
       nclasses = 2
       outputs = np.where(outputs > 0.5, 1, 0)
   else:
       # 1-of-N encoding
       outputs = np.argmax(outputs, 1)
       targets = np.argmax(targets, 1)
   cm = np.zeros((nclasses, nclasses))
   for i in range(nclasses):
       for j in range(nclasses):
           cm[i, j] = np.sum(
              np.where(outputs == i, 1, 0) * np.where(targets == j, 1, 0)
           )
   output = cm
   print("Percentage Correct: ", np.trace(cm) / np.sum(cm) * 100)
   return output
```

Helper functions:-

To display confusion matrix-

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

```
def displayConfusionMatrix(cm, plt):
    out_cm = np.array(cm)
    df_cm = pd.DataFrame(out_cm)
    plt.figure(figsize=(10,7))
    sn.set(font_scale=1) # for label size
    sn.heatmap(df_cm, annot=True, annot_kws={"size": 14}) # font size
To show classification report:-
from sklearn.metrics import classification_report
def printClassificationReport(network, test, testt):
    targets=testt
    inputs = np.concatenate((test, -np.ones((np.shape(test)[0], 1))), axis=1)
    nclasses = np.shape(targets)[1]
    output = network.mlpfwd(inputs)
    if nclasses == 1:
        nclasses = 2
        output = np.where(output > 0.5, 1, 0)
    else:
        # 1-of-N encoding
        output = np.argmax(output, 1)
        targets = np.argmax(targets, 1)
    print(classification_report(targets, output))
Including 'pima-indian-diabetes' dataset:-
   loading data
                                                                                               ▷ ↑ ↓ ■ ··· •
       #loading data
       dataset = np.loadtxt('diabetes.csv',delimiter=',')
       no_of_columns = 8
       #normalizing the input
       dataset[:,:no_of_columns] = dataset[:,:no_of_columns]-dataset[:,:no_of_columns].mean(axis=0)
       imax = np.concatenate((dataset.max(axis=0)
             *np.ones((1,no_of_columns+1)),np.abs(dataset.min(axis=0)
              *np.ones((1,no_of_columns+1)))),axis=0).max(axis=0)
       dataset[:,:no_of_columns] = dataset[:,:no_of_columns]/imax[:no_of_columns]
     V 025
                                                                                                            Python
      print(dataset.shape)
[120] 🗸 0.3s
                                                                                                            Python
 ... (768, 9)
       # Split into training, validation, and test sets
       target = np.zeros((np.shape(dataset)[0],2))
       indices = np.where(dataset[:,no_of_columns]=0)
       target[indices,0] = 1
       indices = np.where(dataset[:,no_of_columns]=1)
       target[indices, 1] = 1
       # Randomly order the data
```

order = np.arange(np.shape(dataset)[0])

```
np.random.shuffle(order)
        dataset = dataset[order,:]
        target = target[order,:]
        train = dataset[::2,0:no_of_columns]
        traint = target[::2]
        valid = dataset[1::4,0:no_of_columns]
         validt = target[1::4]
         test = dataset[3::4,0:no_of_columns]
        testt = target[3::4]
        print (train.max(axis=0), train.min(axis=0))
[121] 🗸 0.4s
                                                                                                                                                           Python

    0.04006288
    0.64965237
    1.
    0.78334653
    1.

    1.
    ] [-0.29228942
    -1.
    -1.

    i9 -0.2563047
    ]
    -1.

               0.64606288 0.64965237 1.
... [1.
      1.
                                                                               -0.26173249 -0.1041496 -0.91127677
      -0.20218239 -0.2563047 ]
```

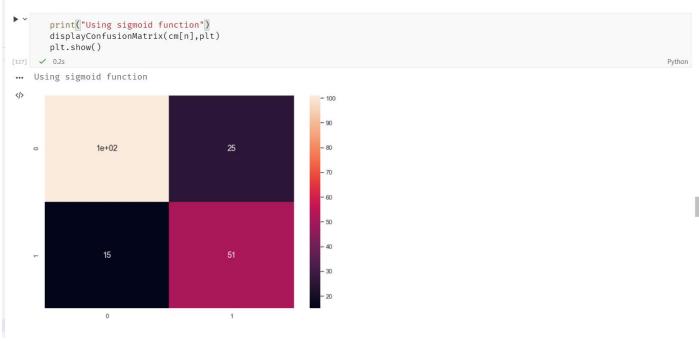
Find the number of neurons

using sigmoid function

```
print("Using Sigmoid function")
       acc = np.zeros(30)
       err = np.zeros(30)
       cm = []
       net = []
       for i in range(30):
           net.append(mlp(train,traint,i+1, outtype='logistic'))
           err[i] = net[i].earlystopping(train,traint,valid,validt,0.001)
           #err = net.mlptrain(train, traint, 0.25, 10000)
           cm.append(net[i].confmat(test,testt))
           acc[i] = np.trace(cm[i]) / np.sum(cm[i]) * 100
[122] 🗸 20.6s
... Using Sigmoid function
    No. of neurons in hidden layers = 1
    Stopped, error = 31.96408393423567
    Percentage Correct: 78.125
    No. of neurons in hidden layers = 2
    Stopped, error = 32.141457877281724
    Percentage Correct: 78.64583333333334
    No. of neurons in hidden layers = 3
    Stopped, error = 32.05194270972359
    Percentage Correct: 78.125
    No. of neurons in hidden layers = 4
    Stopped, error = 32.03907183671007
    Percentage Correct: 77.6041666666666
       plt.xlabel("Number of neurons")
       plt.ylabel("Accuracy")
       plt.plot(acc, color='green')
       plt.show()
[123] 🗸 0.2s
                                                                                                                                 Python
     80.0
                                                                                                                             77.5
     75.0
     72.5
     70.0
     65.0
     62.5
          0
                      10 15 20
Number of neurons
```

```
plt.xlabel("Number of neurons")
plt.ylabel("Error")
        plt.plot(err, color='red')
        plt.show()
[124] 		0.2s
                                                                                                                                                Python
                                                                                                                                           40
    D 36
                 5
          0
                       10 15 20
Number of neurons
         n = np.argmax(acc)
         print("Number of neurons for maximum accuracy =", n)
         print("Accuracy = ", acc[n])
 [125] 🗸 0.3s
  ... Number of neurons for maximum accuracy = 11
      Accuracy = 79.1666666666666
       cm[n]
 [126] 🗸 0.3s
 ... array([[101., 25.], [ 15., 51.]])
```

Using 11 neurons gave the maximum accuracy for sigmoid activation function.



```
print("Classification report using sigmoid activation function")
       print\overline{C}lassificationReport(net[n],\ test,\ testt)
[128] 		 0.3s
... Classification report using sigmoid activation function
                 precision recall f1-score support
              0
                      0.80
                                0.87
                                          0.83
                                                   116
              1
                      0.77
                                0.67
                                          0.72
                                                    76
                                          0.79
                                                    192
       accuracy
                      0.79
                                0.77
                                          0.78
                                                    192
       macro avg
                      0.79
    weighted avg
                                0.79
                                          0.79
                                                    192
```

Trying out different activation functions:-

I. Softmax activation function

softmax activation function

```
print("Using Softmax activation function")
       softnet = mlp(train,traint,n, outtype='softmax')
      softnet.earlystopping(train,traint,valid,validt,0.25)
      softcm = softnet.confmat(test, testt)
[139] 🗸 1.2s
... Using Softmax activation function
    No. of neurons in hidden layers = 11
    Stopped, error = 32.36775940519345
    Percentage Correct: 79.1666666666666
      print("Using softmax function")
      displayConfusionMatrix(softcm,plt)
      plt.show()
[140] 🗸 0.2s
... Using softmax function
</>
                                                                                                                     <u>III</u>
               1e+02
      print("Classification report using Softmax activation function")
      printClassificationReport(softnet, test, testt)
\dots Classification report using Softmax activation function
                 precision recall f1-score support
                                                 116
               0
                      0.79
                            0.89
                                        0.84
                            0.64
                                        0.71
              1
                      0.79
                                                    76
                                         0.79
                                                    192
       accuracy
       macro avg
                      0.79
                              0.77
                                         0.77
                                                    192
                      0.79
                            0.79
                                         0.79
                                                    192
    weighted avg
```

II. Linear activation function



III. Tanh activation function

macro avg

weighted avg

Tanh activation function

0.77

0.77

0.75

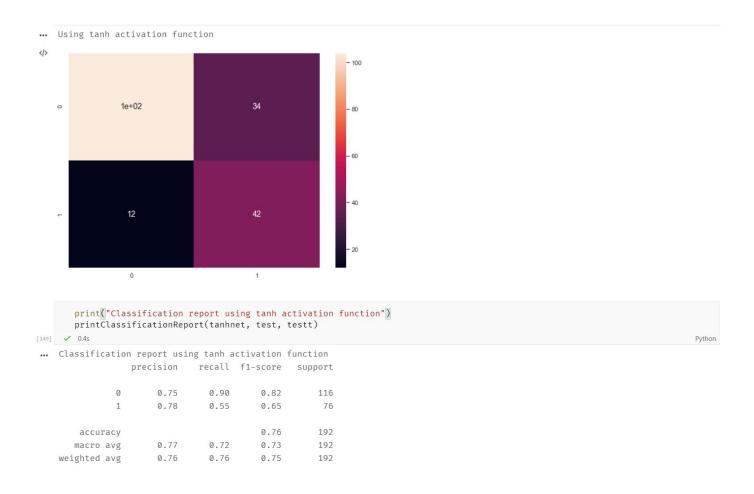
0.78

0.76

0.77

192

192

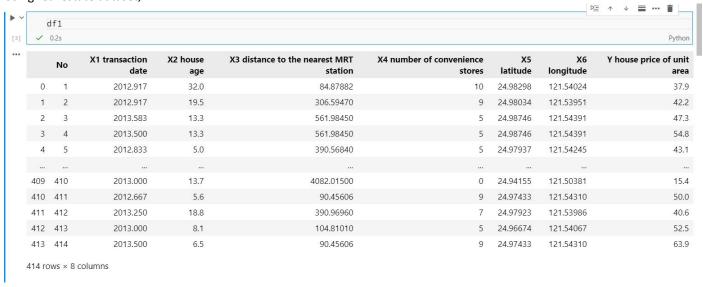


Result:-

For the given dataset, both Sigmoid and Softmax give the highest accuracy of about 79% while linear activation gives 77% accuracy and tanh follows very closely by 76% of test accuracy.

2. Choose a dataset suitable for regression and apply regression using MLP

Using real estate dataset,



```
import numpy as np
      import pandas as pd
      from sklearn.neural_network import MLPRegressor
      from sklearn.metrics import r2_score
      df1 = pd.read_csv('Real estate.csv')
      data1 = df1.drop(['Y house price of unit area'], axis = 1)
      target1 = df1['Y house price of unit area']
      from sklearn.model_selection import train_test_split
      datasets = train_test_split(data1, target1, test_size=0.3)
      X_train, X_test, y_train, y_test = datasets
      reg = MLPRegressor(hidden_layer_sizes=(64,64,64),activation="relu" , max_iter=5000).fit(X_train, y_train)
                                                                                                                                 Python
                                                                                                                □ ↑ ↓ ■ … 1
      y\_pred=reg.predict(X\_test)
      print("The Score with ",(r2_score(y_pred, y_test)))
                                                                                                                                 Python
   The Score with 0.5128951299414292
      import matplotlib.pyplot as plt
      import seaborn as sns
      plt.style.use('ggplot')
      plt.figure(figsize=(10,10))
      sns.regplot(y_test, y_pred, fit_reg=True, scatter_kws={"s": 100})
      plt.show()
[6] 🗸 2.1s
</>
                                                                                                                            60
                            Y house price of unit area
```