# Deep Learning Lab Exam Unsolved Fall 2023

November 8, 2024

### 1 CS 316: Introduction to Deep Learning

#### 2 Lab Exam - Fall 2023

#### 3 Dr. Abdul Samad

Total Duration: 3 Hours Total Points: 100 Name: Write your Name Here ID: Write your Student ID

#### 4 Instructions

- 1. Google Colab must be used for this exam.
- 2. You are not permitted to utilize the internet or any other source for this exam.
- 3. Any violation shall be treated as a plagiarism case.
- 4. The error in one task will not carry to other tasks.
- 5. The marks for each task are stated explicitly.
- 6. Please carefully study the questions; they are self-explanatory.
- 7. Rename your file as Lab\_Exam\_aa01234.ipynb where aa01234 will be replaced by your student id
- 8. The total duration for their exam is 3 hours whereas the last 15 minutes are for submission.

#### 5 Exam Overview

In this Exam, we are implementing a neural network with 3-hidden layers and we will be using it to compute the progression of diabetes. Since each example in the dataset has 10 attributes, the input vector will be of size 10.

The neural network is defined as follows:

$$\mathbf{z}_1 = \mathbf{W}_1 \mathbf{X} + \mathbf{b}_1 \tag{1.1}$$

$$\mathbf{a}_1 = ReLU(\mathbf{z}_1) \tag{1.2}$$

$$\mathbf{z}_2 = \mathbf{W}_2 \mathbf{a}_1 + \mathbf{b}_2 \tag{1.3}$$

$$\mathbf{a}_2 = Tanh(\mathbf{z_2}) \tag{1.4}$$

$$\mathbf{z}_3 = \mathbf{W}_3 \mathbf{a_2} + \mathbf{b}_3 \tag{1.5}$$

$$\mathbf{a}_3 = ReLU(\mathbf{z_3}) \tag{1.6}$$

$$\mathbf{z}_4 = \mathbf{W}_4 \mathbf{a}_3 + \mathbf{b}_4 \tag{1.7}$$

$$\mathbf{a}_4 = \mathbf{z}_4 \tag{1.8}$$

$$\mathbf{\hat{y}} = \mathbf{W}_4 \mathbf{a}_4 + \mathbf{b}_4 \tag{1.}$$

$$L^{(i)} = (\mathbf{y}^{(i)} - \hat{\mathbf{y}}^{(i)})^2 \tag{1.10}$$

$$J = \frac{1}{m} \sum_{i=1}^{m} L^{(i)} \tag{1.11}$$

### 6 Setup

```
[1]: # DO NOT EDIT
     # Import necessary libraries
     # Import the NumPy library for numerical operations.
     import numpy as np
     # Import the Pandas library for data manipulation and analysis.
     import pandas as pd
     # Import the Matplotlib library for data visualization.
     import matplotlib.pyplot as plt
     # Import the Seaborn library for data visualization
     import seaborn as sns
     # Import the datasets module from scikit-learn, a machine learning library.
     from sklearn import datasets
     from sklearn.datasets import load_diabetes
     # Import the StandardScaler for feature scaling.
     from sklearn.preprocessing import StandardScaler
     from sklearn.metrics import r2_score
```

# 7 Diabetes Progression Dataset

Let us import the dataset & create a dataframe

```
[2]: # DO NOT EDIT
    # Load the diabetes dataset
    diabetes = datasets.load_diabetes()
```

Let us learn more about the dataset

```
[3]: # DO NOT EDIT

# Print the description of the dataset

print(diabetes.DESCR)
```

.. \_diabetes\_dataset:

```
Diabetes dataset
```

Ten baseline variables, age, sex, body mass index, average blood

pressure, and six blood serum measurements were obtained for each of n = 442 diabetes patients, as well as the response of interest, a quantitative measure of disease progression one year after baseline.

\*\*Data Set Characteristics:\*\*

:Number of Instances: 442

:Number of Attributes: First 10 columns are numeric predictive values

:Target: Column 11 is a quantitative measure of disease progression one year after baseline

:Attribute Information:

- age age in years
- sex
- bmi body mass index
- bp average blood pressure
- s1 tc, total serum cholesterol
- s2 ldl, low-density lipoproteins
- s3 hdl, high-density lipoproteins
- s4 tch, total cholesterol / HDL
- s5 ltg, possibly log of serum triglycerides level
- s6 glu, blood sugar level

Note: Each of these 10 feature variables have been mean centered and scaled by the standard deviation times the square root of `n\_samples` (i.e. the sum of squares of each column totals 1).

#### Source URL:

https://www4.stat.ncsu.edu/~boos/var.select/diabetes.html

For more information see:

Bradley Efron, Trevor Hastie, Iain Johnstone and Robert Tibshirani (2004) "Least Angle Regression," Annals of Statistics (with discussion), 407-499. (https://web.stanford.edu/~hastie/Papers/LARS/LeastAngle\_2002.pdf)

Checking the Feature Names

- [4]: # DO NOT EDIT
  diabetes.feature\_names
- [4]: ['age', 'sex', 'bmi', 'bp', 's1', 's2', 's3', 's4', 's5', 's6']

Checking the shape of data

[5]: # DO NOT EDIT
diabetes.data.shape

```
[5]: (442, 10)
     There are 10 variables & 442 instances/rows
 [6]: # DO NOT EDIT
      diabetes.target.shape
 [6]: (442,)
     The Target Variable has 442 rows
     Let us Create a Dataframe using this data
 [7]: # DO NOT EDIT
      db df = pd.DataFrame(diabetes.data, columns=diabetes.feature names)
     Display a random sample of 5 rows from the DataFrame
 [8]: # DO NOT EDIT
      db_df.sample(5)
 [8]:
                                    bmi
                                                bp
                                                          s1
                                                                    s2
                                                                               s3
                age
                          sex
      334 -0.060003 0.050680 -0.047163 -0.022885 -0.071743 -0.057681 -0.006584
      77 -0.096328 -0.044642 -0.036385 -0.074527 -0.038720 -0.027618
      436 -0.056370 -0.044642 -0.074108 -0.050427 -0.024960 -0.047034
      376 -0.001882 -0.044642 0.068163 -0.005670 0.119515 0.130208 -0.024993
      241 0.030811 0.050680 -0.008362 0.004658 0.014942 0.027496 0.008142
                 s4
                           s5
                                      s6
      334 -0.039493 -0.062917 -0.054925
      77 -0.039493 -0.074093 -0.001078
      436 -0.076395 -0.061176 -0.046641
      376 0.086708 0.046133 -0.001078
      241 -0.008127 -0.029526 0.056912
     Add Dependent Variable to the dataset
 [9]: # DO NOT EDIT
      db_df['Progression'] = diabetes.target
     Display a random sample of 5 rows from the DataFrame
[10]: # DO NOT EDIT
      db_df.sample(5)
[10]:
                                    bmi
                                                          s1
                                                                    s2
                                                                               s3
                                                                                   \
                age
                          sex
                                                bp
      178 0.041708 -0.044642 -0.008362 -0.026328
                                                    0.024574
                                                              0.016222 0.070730
          -0.089063 -0.044642 -0.011595 -0.036656
                                                    0.012191
                                                              0.024991 -0.036038
      173 -0.063635 0.050680 -0.079497 -0.005670 -0.071743 -0.066449 -0.010266
         -0.016412 -0.044642 -0.035307 -0.026328 0.032830
                                                              0.017162 0.100183
```

0.019913 - 0.044642 - 0.023451 - 0.071085 0.020446 - 0.010082 0.118591

```
    s4
    s5
    s6
    Progression

    178
    -0.039493
    -0.048359
    -0.030072
    81.0

    3
    0.034309
    0.022688
    -0.009362
    206.0

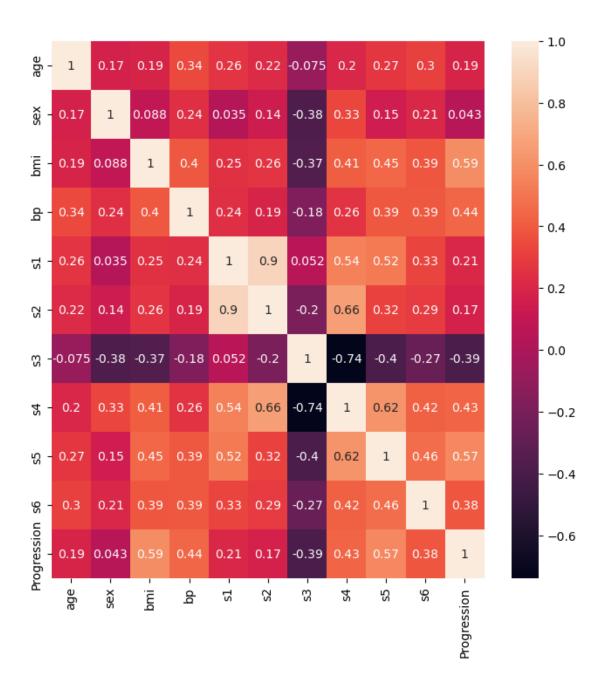
    173
    -0.039493
    -0.018114
    -0.054925
    101.0

    82
    -0.039493
    -0.070209
    -0.079778
    52.0

    43
    -0.076395
    -0.042571
    0.073480
    92.0
```

Let us visualize the Linear correlation between the variables in the dataframe by plotting a heatmap.

```
[11]: # DO NOT EDIT
plt.subplots(figsize=(8,8))
sns.heatmap(db_df.corr(),annot=True)
plt.show()
```



# 8 Data Modelling

```
[12]: # DO NOT EDIT
# Independent variables / explanatory variables
# Axis=1 means we drop data by column.
X = db_df.drop(labels='Progression', axis=1).values
# Dependent variable / response / target variable.
Y = db_df['Progression'].values
```

```
# Reshape the target variable
Y = Y.reshape(len(Y),1)
print(X.shape,Y.shape)
```

(442, 10) (442, 1)

### 9 [10 Points] Task 01 - Train Test Split

In Task 01, you are required to the implement the function train\_test\_split which splits the datasets into train and test set.

Hint: Use rng.shuffle or rng.permutation to shuffle the dataset. Furthermore, instead of shuffling the dataset, shuffle the indexes.

```
[13]: #TODO: Implement the function train_test_split
      def train_test_split(inputs,outputs,test_size,seed = 0):
          Splits the data into training and test sets.
          Return 4 numpy arrays. X_train, X_test, Y_train, Y_test
          where training data is test_size proportion of data provided.
              inputs [np.array] : numpy array of input data
              outputs [np.array]: numpy array of output labels
              test\_size [float]: proportion of data to be used as test data. e.g. 0.2 \Box
       \hookrightarrowmeans 20% of data is used for test data.
              seed [int]: A seed to create random number generator. (For ...
       \neg reproducability)
          n n n
          rng = np.random.default rng(seed)
          assert(len(inputs) == len(outputs))
          assert(test_size <= 1.0)</pre>
          assert(test_size >= 0.0)
          num_samples = len(inputs)
          num_train = int(num_samples * (1.0 - test_size))
          # Write your code here
          indices = np.arange(num_samples)
          rng.shuffle(indices)
          train_indices = indices[:num_train]
          test_indices = indices[num_train:]
          X_train = inputs[train_indices]; X_test = inputs[test_indices]
          Y_train = outputs[train_indices]; Y_test = outputs[test_indices]
          return X_train, X_test, Y_train, Y_test
```

```
[14]: # DO NOT EDIT
      # Test Case to check Train Test Split
      np.random.seed(1)
      x = np.random.randn(15,2)
      y = np.random.randn(15)
      X_train, X_test, Y_train, Y_test = train_test_split(x, y, test_size=0.2)
      print(X_train.shape, X_test.shape, Y_train.shape, Y_test.shape)
      assert np.allclose (X_train,np.array([[ 0.86540763, -2.3015387 ],[ 0.90159072, _
       -0.50249434],[ 1.74481176, -0.7612069 ],[-1.10061918, 1.14472371],[ 1.
       →62434536, -0.61175641],[ 0.3190391 , -0.24937038],[ 1.13376944, -1.
       $\text{\circ}$09989127],[ 1.46210794, -2.06014071],[-0.26788808, 0.53035547],[ 0.
       90085595, -0.68372786],[-0.3224172, -0.38405435],[0.04221375, 0.
       →58281521]]))
      assert np.allclose (X_test,np.array([[-0.12289023, -0.93576943],[-0.17242821,__
       \leftarrow-0.87785842],[-0.52817175, -1.07296862]]))
      assert np.allclose (Y_train,np.array([-0.6871727 , -0.88762896, -0.84520564, -0.
       →19183555, -0.69166075,-0.67124613, 0.2344157 , -0.0126646 , 0.05080775, -0.
       →74715829,-1.11731035, 0.74204416]))
      assert np.allclose (Y test, np.array([ 1.6924546 , 1.65980218, -0.39675353]))
```

(12, 2) (3, 2) (12,) (3,)

The expected output is as follows:

## 10 [10 Points] Task 02 - Activation Functions

In this task, you will be implementing the Tanh and LeakyReLU by completing the Tanh and LeakyReLU classes.

#### 10.1 [5 Points] Part A - LeakyReLU Activation Function

The LeakyReLU activation function and its dervivative are defined as follows:

$$f(z) = \begin{cases} z & z > 0\\ \alpha \cdot z & z \le 0 \end{cases}$$
 (2.1)

$$f'(z) = \begin{cases} 1 & z > 0 \\ \alpha & z \le 0 \end{cases}$$
 (2.2)

```
[15]: #TODO: Implement the LeakyReLU class which implements the LeakyReLU activation

if unction.

class LeakyReLU:

"""

Implements the LeakyReLU activation

"""

def __init__(self, alpha=0.01):
```

#### 10.2 [5 Points] Part B - Tanh Activation Function

The tanh activation function and its derivative are defined as follows:

$$tanh(z) = \frac{e^z - e^{-z}}{e^z + e^{-z}} \tag{3.1}$$

$$tanh'(z) = 1 - tanh(z)^2 \tag{3.2}$$

#### Hint: Use the np.tanh(x) function

```
[20]: #TODO: Implement the Tanh class which implements the Tanh activation function.
      class Tanh:
        11 11 11
        Implements the Tanh activation
        def forward(self, x):
            # Write your code here
            # Hint: Use np.tanh
            # Refer to equation (3.1)
            numer = np.subtract(np.exp(x), np.exp(-x))
            denom = np.add(np.exp(x), np.exp(-x))
            return np.divide(numer, denom)
        def backward(self, x):
            # Write your code here
            # Hint: Use np.tanh
            # Refer to equation (3.2)
            return 1 - np.square(np.tanh(x))
```

```
[22]: # DO NOT EDIT
    # Test Case to Check Tanh
    rng = np.random.default_rng(42)
    input = rng.normal(size=(3,3))
    th = Tanh()
    print(f"Input: {input}")
    output =th.forward(input)
    print(f"Output: {output}")
    assert np.allclose(output,np.array([[ 0.29562343, -0.77788179,  0.63541805],[ 0.47354816 , -0.96039983, -0.86228318],[ 0.12714849, -0.30610551, -0.401679958]]))
    grad = th.backward(input)
    print(f"Gradient: {grad}")
    assert np.allclose(grad,np.array([[0.91260679,  0.39489992,  0.5962439 ],[ 0.45906681,  0.07763216,  0.25646771],[ 0.983833326,  0.90629941,  0.99971777]]))
```

```
Input: [[ 0.30471708 -1.03998411  0.7504512 ]
  [ 0.94056472 -1.95103519 -1.30217951]
```

```
[ 0.1278404 -0.31624259 -0.01680116]]
Output: [[ 0.29562343 -0.77788179  0.63541805]
  [ 0.7354816  -0.96039983 -0.86228318]
  [ 0.12714849 -0.30610551 -0.01679958]]
Gradient: [[0.91260679  0.39489992  0.5962439 ]
  [0.45906681  0.07763216  0.25646771]
  [0.98383326  0.90629941  0.99971777]]
```

### 11 [20 Points ] Task 03 - Dense Layer

In this task, you are provided with a partial implementation of the DenseLayer class. Your are required to complete the class by filling in the missing code blocks.

- 1. Initialization (\_\_init\_\_ method):
- Set the weight matrix (self.weights) using Xavier initialization i.e.

$$W \sim N(0, 2/(input\_dim + output\_dim))$$
(3.1)

where  $input\_dim$  is the number of input neurons and  $output\_dim$  is the number of output neurons.

- Initialize the bias vector (self.bias) with zeros.
- Assign the appropriate activation function based on the given activation parameter.
- 2. Forward Pass (forward method):
- Calculate the linear transformation which is defined as follows:

$$z = \langle XW \rangle + b \tag{3.2}$$

$$a = \sigma(z) \tag{3.3}$$

where W represents self.weights, b represents self.bias, X represents input,  $\sigma$  denotes the activation function, and  $\langle \rangle$  represents dot product (np.dot)

• In case, no activation function is specified:

$$a = z \tag{3.4}$$

- 3. Backward Pass (backward method):
- Compute d\_z,d\_activation, d\_input, d\_weights and d\_bias
- d\_activation represents the derivative of the activation function  $(\delta_{activation})$
- d\_z represents the derivative of the loss function J with respect to the weighted sum before applying an activation function.  $(\delta_z)$

$$\delta_z = \delta_{output} \odot \delta_{activation} \tag{3.5}$$

where  $\odot$  represents elementwise multiplication (hadmard product)

• In case, no activation function was applied:

$$\delta_z = \delta_{output} \tag{3.6}$$

• d\_input represents the derivative of the loss function J with respect to input  $(\delta_{input})$ 

$$\delta_{input} = \langle \delta_z, W.T \rangle \tag{3.7}$$

where W represents self.weights and  $\langle \rangle$  denotes dot product

• d\_weights represents the derivative of the loss function J with respect to weights  $(\delta_{weights})$ 

$$\delta_{weights} = \langle X.T, \delta_z \rangle \tag{3.8}$$

where X represents self.input and  $\langle \rangle$  denotes dot product

• d\_bias represents the derivative of the loss function J with respect to bias  $(\delta_{bias})$ 

$$\delta_{bias} = \sum_{i} \delta_z \tag{3.9}$$

```
[23]: #TODO: Complete the class DenseLayer
      class DenseLayer:
          def __init__(self, input_dim, output_dim, activation):
              # Refer to equation (3.2)
              # Hint: Use np.random.normal
              self.weights = np.random.normal(0, np.sqrt(2/(input_dim + output_dim)),__
       →(input_dim, output_dim))
              # Hint: Use np.zeros
              self.bias = np.zeros((output_dim))
              if activation == 'leakyrelu':
                  self.activation = LeakyReLU()
              elif activation == 'tanh':
                  self.activation = Tanh()
              elif activation == 'linear':
                  # Linear activation (no activation)
                  self.activation = None
              else:
                  raise ValueError("Unsupported activation function")
          def forward(self, x):
              self.input = x
              # Refer to equation (3.2)
              self.z = np.dot(x, self.weights) + self.bias
              if self.activation:
                  # Refer to equation (3.3)
                  self.output = self.activation.forward(self.z)
              else:
                  # No activation (linear)
                  # Refer to equation (3.4)
```

```
self.output = self.z
    return self.output
def backward(self, d_output):
    if self.activation:
        d_activation = self.activation.backward(self.z)
        # Refer to equation (3.5)
        d_z = d_output * d_activation
    else:
        # Refer to equation (3.6)
        d_z = d_output
    # Refer to equation (3.7)
    d_input = np.dot(d_z, self.weights.T)
    # Refer to equation (3.8)
    d_weights = np.dot(self.input.T, d_z)
    # Refer to equation (3.9)
    d_bias = np.sum(d_z, axis=0)
    return d_input, d_weights, d_bias
```

```
[24]: # DO NOT EDIT
      # Test code for checking function
      # Initialise Layer
      D = DenseLayer(3,5,'tanh')
      # Expected output
      print(f'Weights shape {D.weights.shape}\nBias shape {D.bias.shape}\nActivation⊔
       →{D.activation}')
      # Check dimensions
      assert D.weights.shape == (3,5) and D.bias.shape == (5,)
      rng = np.random.default rng(42)
      # Define sample input
      sample_x = rng.normal(3,3,size=(2,3))
      # Compute forward pass
      sample_output = D.forward(sample_x)
      # Expected ouput
      print(f'Input shape {sample_x.shape}\nOutput shape {sample_output.shape}')
      assert sample_output.shape == (2,5)
      # Define sample grad
      sample_grad = rng.normal(3,3,size=(2,5))
      # Compute backward_pass
      sample_d input,sample_d weights,sample_d bias = D.backward(sample grad)
      print(f'Grad Input shape {sample_d_input.shape}\nGrad Output shape_
       sample_d_weights.shape}\nGrad_Bias_shape {sample_d_bias.shape}')
      assert sample_d_input.shape == (2,3)
      assert sample d weights.shape == (3,5)
```

```
assert sample_d_bias.shape == (5,)
```

```
Weights shape (3, 5)
Bias shape (5,)
Activation <__main__.Tanh object at 0x74a1909bfd40>
Input shape (2, 3)
Output shape (2, 5)
Grad Input shape (2, 3)
Grad Output shape (3, 5)
Grad Bias shape (5,)
```

The expected output is as follows. Since the actual values might differ due to randomness, the expected output includes dimensions.

#[10 Points] Task 04 - AdamOptimizer

In this task, you are provided with a partial implementation of the AdamOptimizer class. Your are required to complete the class by filling in the missing code blocks.

In the apply\_gradients method, for each parameter i: - Calculate the fist moment estimate:

$$m_i = \beta_1 \times m_i + (1 - \beta_1) \times grads_i \tag{4.1}$$

- Calculate the second moment estimate:

$$v_i = \beta_2 \times v_i + (1 - \beta_2) \times (grads_i)^2 \tag{4.2}$$

- Compute the bias-corrected first moment estimate:

$$\hat{m}_i = \frac{m_i}{1 - \beta_1^t} \tag{4.3}$$

- Compute the bias-corrected second moment estimate:

$$\hat{v}_i = \frac{v_i}{1 - \beta_2^t} \tag{4.4}$$

- Update the parameter:

$$\theta_{t+1} = \theta_t - \frac{\eta}{\sqrt{\hat{v}_t} + \epsilon} \hat{m}_t \tag{4.5}$$

```
self.m_hat = None
              self.v = None
              self.v hat = None
              self.t = 0
          def apply_gradients(self, params, grads):
              if self.m is None:
                  self.m = [np.zeros(param.shape) for param in params]
                  self.m hat = [np.zeros(param.shape) for param in params]
                  self.v = [np.zeros(param.shape) for param in params]
                  self.v hat = [np.zeros(param.shape) for param in params]
              self.t += 1
              for i in range(len(params)):
                  # Refer to equation (4.1)
                  self.m[i] = (self.beta1 * self.m[i]) + ((1 - self.beta1) * grads[i])
                  # Refer to equation (4.2)
                  self.v[i] = (self.beta2 * self.v[i]) + ((1 -self.beta2) * np.

square(grads[i]))
                  # Refer to equation (4.3)
                  self.m hat[i] = self.m[i]/(1 - self.beta1**self.t)
                  # Refer to equation (4.4)
                  self.v_hat[i] = self.v[i]/(1 - self.beta2**self.t)
                  # Refer to equation (4.5)
                  params[i] -= (self.learning_rate * self.m_hat[i])/(np.sqrt(self.
       →v_hat[i]) + self.epsilon)
[28]: # DO NOT EDIT
      # Test code for checking function
      rng = np.random.default_rng(42)
      test_params = [rng.normal(0,1,size=(i,i)) for i in range(2,4)]
      test params cp = test params[:]
      test_grad = [rng.normal(1,1,size=(i,i)) for i in range(2,4)]
      ad = AdamOptimizer()
      ad.apply_gradients(test_params,test_grad)
      result_params = [np.array([[ 0.30371708, -1.04098411],[ 0.7494512 , 0.
       493956472]]), np.array([[-1.95203519, -1.30317951, 0.1268404],[-0.31724259,
       4-0.01780116, -0.85404393],[ 0.87839797, 0.77679194, 0.0650307 ]])]
      for res,test,grad,i in zip(test_params_cp,result_params,test_grad,range(0,2)):
       print(f'Parameter {i}: {res}\nGradient: {grad}\nResult{test}')
      for res,test in zip(test_params,result_params):
```

assert np.allclose(res,test)

### 12 [10 Points] Task 05 - MultiLayerPerceptron

In this task, you are provided with a partial implementation of the MLP class. Your are required to complete the class by filling in the missing code blocks.

```
[29]: #TODO: Complete the class MLP
      class MLP:
        def __init__(self, input_dim, hidden_units, output_dim):
            self.layers = []
            activations = ['tanh', 'leakyrelu', 'leakyrelu']
            for units, activation in zip(hidden_units, activations):
                # Initialise the Dense Layer object in which the input neurons are
       ⇔equal to input_dim and output_neurons are equal to units
                # Also specify the activation function
                self.layers.append(DenseLayer(input_dim, units, activation))
                # Update input_dim for the next layer
                input_dim = units
            # Initialise the Output Layer object in which the input neurons are equal_
       →to input_dim and output_neurons are equal to output_dim
            # Also specify the activation function as linear
            self.layers.append(DenseLayer(input_dim, output_dim, 'linear'))
        def forward(self, x):
            for layer in self.layers:
                # Apply forward pass on the layer using the forward method
                x = layer.forward(x)
            return x
        def backward(self, d_output):
            weights_list = []
            bias_list = []
            for layer in reversed(self.layers):
                # Apply backwrd pass on the layer using the forward method
                d_output, d_weights, d_bias = layer.backward(d_output)
```

```
weights_list.append(d_weights)
  bias_list.append(d_bias)
return weights_list, bias_list
```

```
[30]: # DO NOT EDIT
      # Test Case for checking function
      # Initialise Multi Layer Perceptron
      m = MLP(3,[8, 4, 2],1)
      # Define Weights shape
      weights_shape = [(3,8),(8,4),(4,2),(2,1)]
      # Define Bias shape
      bias_shape = [(8,),(4,),(2,),(1,)]
      # Expected output
      for l,w_shape,b_shape in zip(m.layers,weights_shape,bias_shape):
       print(f'Weights Shape: {l.weights.shape}\nBias Shape: {l.bias.
       ⇔shape}\nActivation: {l.activation}')
      # Checking dimensions are appropriate
      for l,w_shape,b_shape in zip(m.layers,weights_shape,bias_shape):
       assert w_shape == 1.weights.shape
       assert b_shape == l.bias.shape
      # Define sample input
      sample_input = rng.normal(3,3,size=(5,3))
      # Apply forward pass
      sample_result = m.forward(sample_input)
      # Expected output
      for inp,res in zip(sample_input,sample_result):
       print(f'Input Shape: {inp.shape}\nResult Shape: {res.shape}')
      # Checking dimensions are appropriate
      assert len(sample result) == 5
      # Define sample_gradient
      sample_grad = rng.normal(3,3,size=(5,1))
      # Compute Backward Pass
      weights_grad_list, bias_grad_list = m.backward(sample_grad)
      # Expected Output
      for grad_w,grad_b in zip(weights_grad_list,bias_grad_list):
      print(f'Grad Weights Shape: {grad w.shape}\nGrad Bias Shape: {grad b.shape}')
      # Checking dimensions are appropriate
      for grad_w,grad_b,w_shape,b_shape in_

¬zip(weights_grad_list,bias_grad_list,weights_shape[::-1],bias_shape[::-1]):
        print(grad_w.shape,grad_b.shape,w_shape,b_shape)
        assert grad w.shape == w shape
```

```
Weights Shape: (3, 8)
Bias Shape: (8,)
Activation: <__main__.Tanh object at 0x74a19078c920>
Weights Shape: (8, 4)
Bias Shape: (4,)
Activation: <__main__.LeakyReLU object at 0x74a19078e480>
Weights Shape: (4, 2)
Bias Shape: (2,)
Activation: <__main__.LeakyReLU object at 0x74a19078d370>
Weights Shape: (2, 1)
Bias Shape: (1,)
Activation: None
Input Shape: (3,)
Result Shape: (1,)
Grad Weights Shape: (2, 1)
Grad Bias Shape: (1,)
Grad Weights Shape: (4, 2)
Grad Bias Shape: (2,)
Grad Weights Shape: (8, 4)
Grad Bias Shape: (4,)
Grad Weights Shape: (3, 8)
Grad Bias Shape: (8,)
(2, 1) (1,) (2, 1) (1,)
(4, 2) (2,) (4, 2) (2,)
(8, 4) (4,) (8, 4) (4,)
(3, 8) (8,) (3, 8) (8,)
```

assert grad\_b.shape == b\_shape

The expected output is as follows. Since the actual values might differ due to randomness, the expected output includes dimensions.

### 13 [10 Points ] Task 06 - Create Minibatches

In this task, you have to implement a function named create\_minibatches that splits the feature matrix x and the target variable y into batches of size equal to batch\_size.

```
[31]: #TODO: Complete create_minibatches
def create_minibatches(x,y,batch_size):
   for i in range(0, x.shape[0], batch_size):
        x_batch = x[i: i + batch_size]
        y_batch = y[i: i + batch_size]
        yield x_batch,y_batch
```

```
[32]: # DO NOT EDIT

# Test case to check create_mini_batches

rng = np.random.default_rng(42)
  input = rng.random((442,10))
  output = np.arange(442)
  batch = 1

for xbatch,ybatch in create_minibatches(input,output,64):
    print(f"Batch: {batch} xbatch.shape: {xbatch.shape}, ybatch.shape: {ybatch.shape}")
    if batch != 7:
        assert xbatch.shape == (64,10) and ybatch.shape == (64,)
        else:
        assert xbatch.shape == (58,10) and ybatch.shape == (58,)
        batch+=1
```

```
Batch: 1 xbatch.shape: (64, 10), ybatch.shape: (64,)
Batch: 2 xbatch.shape: (64, 10), ybatch.shape: (64,)
Batch: 3 xbatch.shape: (64, 10), ybatch.shape: (64,)
Batch: 4 xbatch.shape: (64, 10), ybatch.shape: (64,)
Batch: 5 xbatch.shape: (64, 10), ybatch.shape: (64,)
Batch: 6 xbatch.shape: (64, 10), ybatch.shape: (64,)
Batch: 7 xbatch.shape: (58, 10), ybatch.shape: (58,)
```

## 14 [10 Points] Task 07 - MSE

The Mean Squared Error is defined as following:

$$L^{(i)} = (\mathbf{y}^{(i)} - \mathbf{\hat{y}}^{(i)})^2 \tag{7.1}$$

$$J = \frac{1}{m} \sum_{i=1}^{m} L^{(i)} \tag{7.2}$$

where m is the number of examples.

In this task, you have to implement a function named mean\_squared\_error that computes the Mean Squared Error (MSE) between two arrays, y\_hat and y where y\_hat is the predicted value and y is the actual value.

```
[33]: #TODO: Complete the function mean_squared_error

def mean_squared_error(y_hat, y):
    # Write your code here
    # Hint: Use np.mean
    # Refer to equations (7.1) & (7.2)
    return np.mean(np.square(y_hat - y))
```

```
[34]: # DO NOT EDIT
    # Test code for checking function
    y_hat = np.arange(5)
    y = np.arange(5,10)
    mse_result = mean_squared_error(y_hat,y)
    print(f'y_hat {y_hat}, y {y}, result: {mse_result}')
    assert mse_result == 25.0
```

```
y_hat [0 1 2 3 4], y [5 6 7 8 9], result: 25.0
```

### 15 [10 Points ] Task 08 - MSE Grad

The derivative of the Mean Squared Error with respected to the predicted value is as follows:

$$\frac{\delta J}{\delta \hat{y}} = 2(\hat{y} - y) \tag{8.1}$$

In this task, you have to implement a function named mean\_squared\_error\_gradient that computes the derivative of Mean Squared Error (MSE) between two arrays, y\_hat and y where y\_hat is the predicted value and y is the actual value.

```
[35]: #TODO: Complete the function mean_squared_error_gradient

def mean_squared_error_gradient(y_hat,y):
    # Write your code here
    # Refer to equations (8.1)
    return 2*(y_hat - y)
```

```
[36]: # DO NOT EDIT
    # Test code for checking function
    y_hat = np.arange(5)
    y = np.arange(5,10)
    mse_grad = mean_squared_error_gradient(y_hat,y)
    print(f'y_hat {y_hat}, y {y}, result: {mse_grad}')
    assert np.array_equal(mse_grad,np.array([-10 ,-10 ,-10 ,-10]))
```

```
y_hat [0 1 2 3 4], y [5 6 7 8 9], result: [-10 -10 -10 -10]
```

The expected output is as follows:

### 16 [10 Points] Task 09 - R2Score

The R2 score is computed as follows:

$$R^{2} = 1 - \frac{\text{sum squared regression (SSR)}}{\text{total sum of squares (SST)}},$$
(9.1)

$$= 1 - \frac{\sum (y_i - \hat{y_i})^2}{\sum (y_i - \bar{y})^2}.$$
 (9.2)

where y is the actual value,  $\hat{y}$  is the predicted value and  $\bar{y}$  is the mean of actual value.

In this task, you have to implement a function named r2\_score that computes the R2Score between two arrays, y\_hat and y where y\_hat is the predicted value and y is the actual value.

```
[37]: #TODO: Complete the function r2_score
def r2_score(y_hat, y):
    #Write your code here
    #Refer to equations (9.1) & (9.2)

# Calculate the mean of y_true
mean_y_true = np.mean(y)

# Calculate SSR (Sum of Squared Residuals)
ssr = np.sum(np.square(y - y_hat))

# Calculate SST (Total Sum of Squares)
sst = np.sum(np.square(y - mean_y_true))

# Calculate R2 score
r2 = 1 - (ssr/sst)
return r2
```

```
[38]: # DO NOT EDIT
    # Test code for checking function
    y_hat = np.arange(5)
    y = np.arange(5,10)
    r2_result = r2_score(y_hat,y)
    print(f'y_hat {y_hat}, y {y}, result: {r2_result}')
    assert r2_result == -11.5
```

y\_hat [0 1 2 3 4], y [5 6 7 8 9], result: -11.5

The expected output is as follows:

### 17 [10 Points ] Task 10 - Fit Function

In this task, you are required to implement the fit function which implements the main training loop for the model.

```
[39]: #TODO: Complete the function which implements the training loop for the model.
      def fit (X,Y,epochs=200, batch_size=64, learning_rate=0.3):
        # Split the dataset into Train and Test. test_size = 0.2 and seed=42
        X_train, X_test, Y_train, Y_test = train_test_split(X,Y,0.2,42)
        # Standardize the features
        scaler = StandardScaler()
        X_train = scaler.fit_transform(X_train)
        X_test = scaler.transform(X_test)
        # Define the model
        input_dim = X_train.shape[1]
        hidden units = [128, 64, 32]
        output_dim = 1
        hidden_units = [128, 64, 32]
        output_dim = 1
        model = MLP(input_dim, hidden_units, output_dim)
        optimizer = AdamOptimizer(learning_rate=0.001)
        for epoch in range(epochs):
          for x_batch,y_batch in create minibatches(X_train,Y_train,batch_size):
            # Compute forward pass on x_batch using the forward method of model
            y_pred = model.forward(x_batch)
            # Compute loss using the mean squared error function
            loss = mean_squared_error(y_pred, y_batch)
            # Compute derivative of the loss using the mean_squared_error_gradient_
       ⇔ function
            d_loss = mean_squared_error_gradient(y_pred, y_batch)
            # Compute backward pass using the backward method of the model by passing_
       \hookrightarrow d loss as a parameter
            d_weights, d_bias = model.backward(d_loss)
            params = [layer.weights for layer in model.layers] + [layer.bias for_
       →layer in model.layers]
```

```
grads = d_weights[::-1] + d_bias[::-1]
    optimizer.apply_gradients(params, grads)

if epoch % 10 == 0:
    print(f'Epoch {epoch}/{epochs}, Loss: {loss}')

# Comput the model prediction on the X_test by using the forward method_
provided by model

y_pred = model.forward(X_test)

# Compute test mse using the mean_squared_error_function, y_pred and Y_test
mse = mean_squared_error(y_pred, Y_test)
print(f'Mean Squared Error (MSE): {mse}')

# Compute test mse using the mean_squared_error_function, y_pred and Y_test
r2 = r2_score(y_pred, Y_test)
print(f'R-Squared (R^2) Score: {r2:.2f}')

fit(X,Y)
```

```
Epoch 0/200, Loss: 24603.382337311665
Epoch 10/200, Loss: 16147.465676081818
Epoch 20/200, Loss: 3949.296830336809
Epoch 30/200, Loss: 2839.972624640326
Epoch 40/200, Loss: 2287.8900373968895
Epoch 50/200, Loss: 2217.0852539241787
Epoch 60/200, Loss: 2190.3262857338673
Epoch 70/200, Loss: 2152.435324980979
Epoch 80/200, Loss: 2110.0572906644366
Epoch 90/200, Loss: 2069.843686522083
Epoch 100/200, Loss: 2033.2787268550865
Epoch 110/200, Loss: 2004.411379064674
Epoch 120/200, Loss: 1975.2454819073087
Epoch 130/200, Loss: 1946.0873067345092
Epoch 140/200, Loss: 1915.1928502140554
Epoch 150/200, Loss: 1881.7118220096852
Epoch 160/200, Loss: 1851.3329845857954
Epoch 170/200, Loss: 1817.3920137905466
Epoch 180/200, Loss: 1788.8850880183065
Epoch 190/200, Loss: 1761.3692463744328
Mean Squared Error (MSE): 3292.2589701010847
R-Squared (R^2) Score: 0.48
```

The expected output is as follows. However, your output might sligtly differ due to randomness.