Homework 4 Spring 2023

Due Date - 04/19/2023, 11:59PM

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### **UNI: Ih3057**

### PART-1: Neural Network from the scratch

For this part, you are not allowed to use any library other than numpy.

In this part, you will will implement the forward pass and backward pass (i.e. the derivates of each parameter wrt to the loss) with the network image uploaded

The weight matrix for the hidden layer is W1 and has bias b1.

The weight matrix for the ouput layer is W2 and has bias b2.

Activatation function is sigmoid for both hidden and output layer

Loss function is the MSE loss

Refer to the below dictionary for dimensions for each matrix

```
In [2]: import numpy as np
   import matplotlib.pyplot as plt
   import pprint
   pp = pprint.PrettyPrinter(indent=4)
   import warnings
   warnings.filterwarnings("ignore")
```

```
In [57]: np.random.seed(0) # don't change this

weights = {
    'W1': np.random.randn(3, 2),
    'b1': np.zeros(3),
    'w2': np.random.randn(3),
    'b2': 0,
}

X = np.random.rand(1000,2)
Y = np.random.randint(low=0, high=2, size=(1000,))
```

```
In [49]: #Sigmoid Function
def sigmoid(z):
    return 1/(1 + np.exp(-z))

In [52]: def forward_propagation(X, weights):
    # Z1 -> output of the hidden layer before applying activation
    # H -> output of the hidden layer after applying activation
    # Z2 -> output of the final layer before applying activation
    # Y -> output of the final layer after applying activation

Z1 = np.dot(X, weights['W1'].T) + weights['b1']
    H = sigmoid(Z1)

Z2 = np.dot(H, weights['W2'].T) + weights['b2']
    Y = sigmoid(Z2)

return Y, Z2, H, Z1
```

```
In [58]: def back_propagation(X, Y_T, weights):
             N = X.shape[0]
             # forward propagation
             Y, Z2, H, Z1 = forward propagation(X, weights)
             L = (1/(2*N_points)) * np.sum(np.square(Y - Y_T))
             # back propagation
             dLdY = 1/N_points * (Y - Y_T)
             dLdZ2 = np.multiply(dLdY, (sigmoid(Z2)*(1-sigmoid(Z2))))
             dLdW2 = np.dot(H.T, dLdZ2)
             dLdb2 = np.sum(dLdZ2)
             dLdH = np.dot(dLdZ2, weights['W2'])
             dHdZ1 = sigmoid(Z1) * (1 - sigmoid(Z1))
             dLdZ1 = np.multiply(dLdH, dHdZ1)
             dLdW1 = np.dot(dLdZ1.T, X)
             dLdb1 = np.sum(dLdZ1, axis=0)
             gradients = {
                 'W1': dLdW1,
                 'b1': dLdb1,
                 'W2': dLdW2,
                 'b2': dLdb2,
             }
             return gradients, L
```

```
In [60]: def back_propagation(X, Y_T, weights):
             N points = X.shape[0]
             # forward propagation
             Y, Z2, H, Z1 = forward propagation(X, weights)
             L = (1/(2*N_points)) * np.sum(np.square(Y - Y_T))
             # back propagation
             dLdY = 1/N_points * (Y - Y_T)
             dLdZ2 = np.multiply(dLdY, (sigmoid(Z2)*(1-sigmoid(Z2))))
             dLdW2 = np.dot(H.T, dLdZ2.reshape(N points, 1))
             dLdb2 = np.sum(dLdZ2, axis=0)
             dLdH = np.dot(dLdZ2.reshape(N points, 1), weights['W2'].reshape(1, 3)
             dLdZ1 = np.multiply(dLdH, (sigmoid(Z1)*(1-sigmoid(Z1))))
             dLdW1 = np.dot(X.T, dLdZ1)
             dLdb1 = np.sum(dLdZ1, axis=0)
             gradients = {
                 'W1': dLdW1,
                 'b1': dLdb1,
                 'W2': dLdW2,
                 'b2': dLdb2,
             }
             return gradients, L
In [61]: gradients, L = back propagation(X, Y, weights)
         print(L)
         0.1332476222330792
In [62]: pp.pprint(gradients)
             'W1': array([[ 0.00244596, -0.00030765, -0.00034768],
                [0.00262019, -0.00024188, -0.000372]
             'W2': array([[0.02216011],
                [0.02433097],
                [0.01797174]]),
             'b1': array([ 0.00492577, -0.00058023, -0.00065977]),
             'b2': 0.029249230265318685}
```

Your answers should be close to L = 0.133 and 'b1': array([ 0.00492, -0.000581, -0.00066])

# **PART 2 MNIST Dataset**

Description: The MNIST dataset is a widely-used benchmark dataset in the field of machine learning and computer vision. It consists of 70,000 grayscale images of handwritten digits (0-9), with 60,000 images in the training set and 10,000 images in the test set. The images are 28x28 pixels in size, and each pixel is represented by an integer value between 0 and 255, with 0 representing a white pixel and 255 representing a black pixel.

```
In [3]: from tensorflow.keras.datasets import mnist

# The MNIST dataset and the labels have been provided for you
(x_dev, y_dev), (x_test, y_test) = mnist.load_data()
In [4]: LABELS = ['0', '1', '2', '3', '4', '5', '6', '7', '8', '9']
```

2.1 Plot 5 samples from each class/label from train set on a 10\*5 subplot

```
In [10]: #Your code here
         import random
         fig,axs = plt.subplots(10,5,figsize=(20,20))
         fig.tight_layout(pad=3)
         for i in range(10):
             target_list = np.where(y_dev==i)[0]
             index = random.choices(list(target_list), k=5)
             x_dev_list = x_dev[index]
             for j in range(5):
                 image = x_dev_list[j]
                 axs[i,j].imshow(image)
                 axs[i,j].set_title(LABELS[i])
```

2.2 Preparing the dataset

- 1) Print the shapes  $x_{dev}$ ,  $y_{dev}$ ,  $x_{test}$ ,  $y_{test}$
- 2) Flatten the images into one-dimensional vectors and again print the shapes of  $x_{dev}$ ,  $x_{test}$
- 3) Standardize the development and test sets.
- 4) Train-test split your development set into train and validation sets (8:2 ratio).

```
In [11]: #Your code here
         print(f"The shape of x_dev is {x_dev.shape}")
         print(f"The shape of y dev is {y dev.shape}")
         print(f"The shape of x test is {x test.shape}")
         print(f"The shape of y_test is {y_test.shape}")
         The shape of x_{dev} is (60000, 28, 28)
         The shape of y_dev is (60000,)
         The shape of x test is (10000, 28, 28)
         The shape of y_test is (10000,)
In [12]: x_{dev_rs} = x_{dev_reshape}(x_{dev_shape}[0], -1)
         x_{test_rs} = x_{test.reshape}(x_{test.shape}[0], -1)
         print(f"The shape of x dev is {x dev rs.shape}")
         print(f"The shape of x_test is {x_test_rs.shape}")
         The shape of x_{dev} is (60000, 784)
         The shape of x test is (10000, 784)
In [13]: from sklearn.preprocessing import StandardScaler
         ss = StandardScaler()
         x_dev_std = ss.fit_transform(x_dev_rs)
         x test std = ss.fit transform(x test rs)
In [14]: from sklearn.model selection import train test split
         from keras.utils.np utils import to categorical
         y dev tc = to categorical(y dev,10)
         y test tc = to categorical(y test, 10)
         x_train, x_val, y_train, y_val = train_test_split(x dev std,y dev tc,test
         Using TensorFlow backend.
         2.3 Build the feed forward network
         First hidden layer size - 128
         Second hidden layer size - 64
```

Third and last layer size - You should know this

```
In [20]: #Your code here
    from tensorflow.keras.models import Sequential
    from tensorflow.keras.layers import Dense, Activation
    from tensorflow.keras import layers
    model = Sequential([
        layers.Dense(128, input_shape=(784,), activation = 'relu'),
        layers.Dense(64, activation = 'relu'),
        layers.Dense(10, activation = 'softmax'),
    ])
```

2.3.1) Comment briefly on importance of activation functions used.

It produces a probability distribution over the classes, which allows the model to make probabilistic predictions for each class. This is useful because it provides a measure of confidence in the model's predictions, and can be used to estimate the uncertainty in the model's predictions.

### 2.4) Print out the model summary

```
In [22]: #Your code here
model.summary()
```

Model: "sequential"

Layer (type)	Output Shape	Param #
dense (Dense)	(None, 128)	100480
dense_1 (Dense)	(None, 64)	8256
dense_2 (Dense)	(None, 10)	650
Total params: 109,386		

Trainable params: 109,386
Non-trainable params: 0

2.5) Do you think this number is dependent on the image height and width?

Yes, I do. If the image height and width were to change, the number of input features would also change accordingly. For example, if the images were resized to 32 by 32 pixels, the number of input features would be 1024. The number of classes is represented in the last layer. The distinct items can be seen in the images. So the number is greater if the pattern is complex.

2.6) Use the right metric and the right loss function and batch size, with Adam as the optimizer, train your model for 10 epochs .

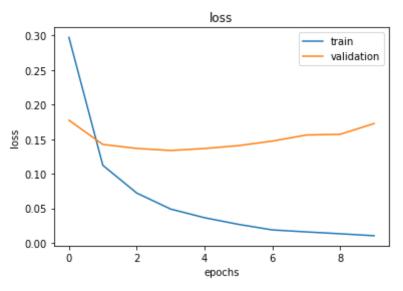
```
In [23]: #Your code here
model.compile(optimizer="adam",loss="categorical_crossentropy",metrics=["
history_callback = model.fit(x_train,y_train,batch_size=128,epochs=10,val
```

```
Train on 48000 samples, validate on 12000 samples
Epoch 1/10
0.2972 - accuracy: 0.9115 - val loss: 0.1774 - val accuracy: 0.9514
Epoch 2/10
0.1122 - accuracy: 0.9660 - val_loss: 0.1424 - val_accuracy: 0.9621
Epoch 3/10
0.0721 - accuracy: 0.9779 - val_loss: 0.1366 - val_accuracy: 0.9650
Epoch 4/10
0.0489 - accuracy: 0.9851 - val loss: 0.1337 - val accuracy: 0.9687
Epoch 5/10
0.0364 - accuracy: 0.9891 - val loss: 0.1365 - val accuracy: 0.9688
Epoch 6/10
0.0268 - accuracy: 0.9922 - val loss: 0.1407 - val accuracy: 0.9693
Epoch 7/10
0.0188 - accuracy: 0.9945 - val loss: 0.1472 - val accuracy: 0.9705
Epoch 8/10
0.0159 - accuracy: 0.9954 - val loss: 0.1560 - val accuracy: 0.9697
Epoch 9/10
0.0132 - accuracy: 0.9961 - val loss: 0.1571 - val accuracy: 0.9705
Epoch 10/10
0.0103 - accuracy: 0.9970 - val loss: 0.1726 - val accuracy: 0.9688
```

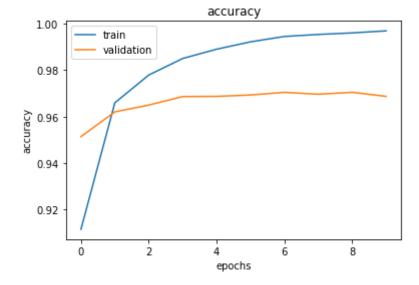
- 2.7) Plot a separate plots for:
- a. displaying train vs validation loss over each epoch
- b. displaying train vs validation accuracy over each epoch

### In [24]: import pandas as pd

```
In [25]: hist_cnn=pd.DataFrame(history_callback.history)
    plt.plot(hist_cnn.index,hist_cnn["loss"])
    plt.plot(hist_cnn.index,hist_cnn["val_loss"])
    plt.xlabel("epochs")
    plt.ylabel("loss")
    plt.title("loss")
    plt.legend(["train","validation"])
    plt.show()
```



```
In [26]: plt.plot(hist_cnn.index,hist_cnn["accuracy"])
    plt.plot(hist_cnn.index,hist_cnn["val_accuracy"])
    plt.xlabel("epochs")
    plt.ylabel("accuracy")
    plt.title("accuracy")
    plt.legend(["train","validation"])
    plt.show()
```



2.8) Finally, report the metric chosen on test set

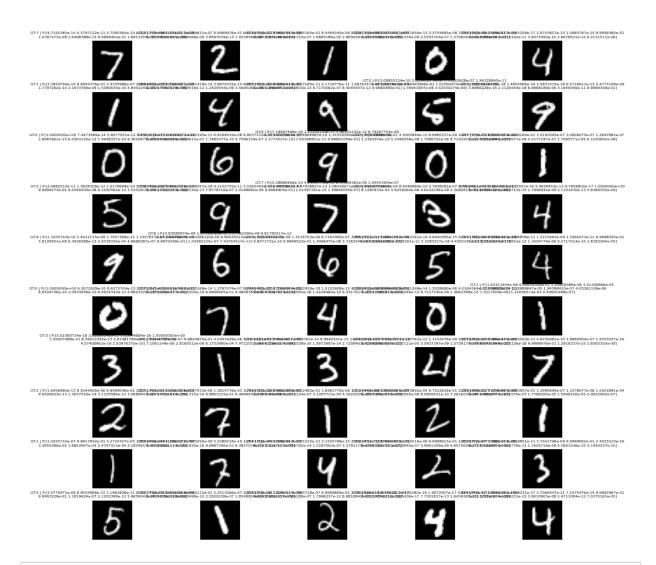
```
In [27]: #Your code here
    score = model.evaluate(x_test_std,y_test_tc,verbose=0)
    print("Test loss: {:.3f}".format(score[0]))
    print("Test accuracy: {:.3f}".format(score[1]))

Test loss: 0.156
    Test accuracy: 0.969
```

2.9 Plot the first 50 samples of test dataset on a 10\*5 subplot and this time label the images with both the ground truth (GT) and predicted class (P).

```
In [40]: import matplotlib.pyplot as plt
         # Get the predicted classes for the test set
         y pred = model.predict(x_test_std[:50])
         # Plot the first 50 samples of the test set
         fig, ax = plt.subplots(10, 5, figsize=(15, 15))
         fig.suptitle('Ground Truth (GT) vs. Predicted Class (P)', fontsize=20)
         for i in range(10):
             for j in range(5):
                 idx = i*5 + j
                 ax[i, j].imshow(x_test[idx], cmap='gray')
                 ax[i, j].set title(f"GT:{y test[idx]} | P:{y pred[idx]}", fontsiz
                 ax[i, j].axis('off')
         # Adjust the spacing between subplots
         plt.subplots_adjust(wspace=0.3, hspace=0.3)
         # Show the plot
         plt.show()
```

#### Ground Truth (GT) vs. Predicted Class (P)



In [ ]: