ICP 4

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GitHub Link: https://github.com/venkat137222/week-4---ICP4

Video Link:

https://drive.google.com/file/d/1sdCV_lznPPaodR5g0mqdDgVHWwjF HCma/view?usp=drive_link

- 1. Use the use case in the class:
- a. Add more Dense layers to the existing code and check how the accuracy changes.

```
import keras
    import pandas
    from keras.models import Sequential
    from keras.layers import Dense, Activation
    from sklearn.model_selection import train_test_split
    import pandas as pd
    import numpy as np
    dataset = pd.read_csv(path_to_csv, header=None).values
    X_train, X_test, Y_train, Y_test = train_test_split(dataset[:,0:8], dataset[:,8],
                                                       test_size=0.25, random_state=87)
    np.random.seed(155)
    my_first_nn = Sequential() # create model
    my_first_nn.add(Dense(20, input_dim=8, activation='relu')) # hidden layer 1
                                                   # Hidden layer 2
    my_first_nn.add(Dense(15, activation='relu'))
                                                             # Hidden layer 3
    my_first_nn.add(Dense(10, activation='relu'))
    my_first_nn.add(Dense(5, activation='relu'))
    my_first_nn.add(Dense(1, activation='sigmoid')) # output layer
    my_first_nn.compile(loss='binary_crossentropy', optimizer='adam', metrics=['acc'])
    my_first_nn_fitted = my_first_nn.fit(X_train, Y_train, epochs=100,
                                        initial epoch=0)
    print(my_first_nn.summary())
    print(my_first_nn.evaluate(X_test, Y_test))
```

Layer (type)	Output Shape	Param #
dense (Dense)	(None, 20)	180
dense_1 (Dense)	(None, 15)	315
dense_2 (Dense)	(None, 10)	160
dense_3 (Dense)	(None, 5)	55
dense_4 (Dense)	(None, 1)	6

```
Total params: 2,150 (8.40 KB)
Trainable params: 716 (2.80 KB)
Non-trainable params: 0 (0.00 B)
Optimizer params: 1,434 (5.61 KB)
None
6/6 ______ 0s 18ms/step - acc: 0.6887 - loss: 0.5574
[0.5714445114135742, 0.6822916865348816]
```

2. Change the data source to Breast Cancer dataset available in the source code folder and make required changes. Report accuracy of the model.

```
# 2. Change the data source to Breast Cancer dataset * available in the source code folder and make required changes.
        # Report accuracy of the model.
[ ] path_to_csv = '/content/gdrive/My Drive/breastcancer.csv
import keras
    import pandas as pd
    import numpy as np
    from keras.models import Sequential
    from keras.layers import Dense, Activation
    from sklearn.model_selection import train_test_split
    dataset = pd.read_csv(path_to_csv)
    X = dataset.loc[:, 'radius_mean':'fractal_dimension_worst']
    Y = dataset['diagnosis']
    # Mapping 'M' to 0 and 'B' to 1 for binary classification
    Y = Y.map({'M': 0, 'B': 1}).astype(int)
    X_train, X_test, Y_train, Y_test = train_test_split(X, Y, test_size=0.25, random_state=87)
    np.random.seed(155)
    my_second_nn = Sequential()
    my_second_nn.add(Dense(20, input_dim=30, activation='relu')) # hidden layer 1
    my_second_nn.add(Dense(10, activation='relu'))
    my_second_nn.add(Dense(5, activation='relu'))
                                                          # Hidden layer 4
    my second nn.add(Dense(1, activation='sigmoid')) # output layer
    my_second_nn.compile(loss='binary_crossentropy', optimizer='adam', metrics=['acc'])
    my_second_nn_fitted = my_second_nn.fit(X_train, Y_train, epochs=100,initial_epoch=0)
    print(my second nn.summary())
    print(my_second_nn.evaluate(X_test, Y_test))
```

```
EPOCH 20/ 100
14/14
                          - 0s 4ms/step - acc: 0.9328 - loss: 0.2499
Epoch 99/100
14/14
                           • 0s 4ms/step - acc: 0.9332 - loss: 0.2461
Epoch 100/100
                          - 0s 5ms/step - acc: 0.9423 - loss: 0.2583
14/14
Model: "sequential_1"
  Layer (type)
                                          Output Shape
                                                                                  Param #
  dense 5 (Dense)
                                           (None, 20)
                                                                                      620
  dense_6 (Dense)
                                           (None, 15)
                                                                                      315
  dense_7 (Dense)
                                           (None, 10)
                                                                                      160
  dense 8 (Dense)
                                           (None, 5)
                                                                                       55
  dense 9 (Dense)
                                           (None, 1)
 Total params: 3,470 (13.56 KB)
 Trainable params: 1,156 (4.52 KB)
```

6

Non-trainable params: 0 (0.00 B) Optimizer params: 2,314 (9.04 KB) None **- 0s** 8ms/step - acc: 0.8775 - loss: 0.3952 [0.3371555209159851, 0.9020978808403015]

3. Normalize the data before feeding the data to the model and check how the normalization change your accuracy (code given below).

from sklearn.preprocessing import StandardScaler sc = StandardScaler()

```
[ ] import keras
    import pandas as pd
    import numpy as np
    from keras.models import Sequential
    from keras.layers import Dense, Activation
    from sklearn.model selection import train test split
    from sklearn.preprocessing import StandardScaler
    dataset = pd.read_csv(path_to_csv)
    X = dataset.loc[:, 'radius_mean':'fractal_dimension_worst']
    Y = dataset['diagnosis']
    # Map 'M' to 0 and 'B' to 1 for binary classification
    Y = Y.map({'M': 0, 'B': 1}).astype(int)
    X_train, X_test, Y_train, Y_test = train_test_split(X, Y, test_size=0.25, random_state=87)
    np.random.seed(155)
    sc = StandardScaler()
    normalized Xtrain = sc.fit transform(X train)
    normalized_Xtest = sc.transform(X_test)
    my third nn = Sequential()
    my_third_nn.add(Dense(20, input_dim=30, activation='relu')) # hidden layer 1
    my_third_nn.add(Dense(15, activation='relu')) # Hidden layer 2
    my_third_nn.add(Dense(10, activation='relu'))
                                                             # Hidden layer 3
    my_third_nn.add(Dense(5, activation='relu'))
                                                             # Hidden layer 4
    my_third_nn.add(Dense(1, activation='sigmoid'))
    my_third_nn.compile(loss='binary_crossentropy', optimizer='adam', metrics=['acc'])
    my_third_nn_fitted = my_third_nn.fit(normalized_Xtrain, Y_train, epochs=100,initial_epoch=0)
    print(my_third_nn.summary())
   print(my_third_nn.evaluate(normalized_Xtest, Y_test))
```

Layer (type)	Output Shape	Param #
dense_10 (Dense)	(None, 20)	620
dense_11 (Dense)	(None, 15)	315
dense_12 (Dense)	(None, 10)	160
dense_13 (Dense)	(None, 5)	55
dense_14 (Dense)	(None, 1)	6

```
Total params: 3,470 (13.56 KB)
Trainable params: 1,156 (4.52 KB)
Non-trainable params: 0 (0.00 B)
Optimizer params: 2,314 (9.04 KB)
None
5/5 _______ 0s 13ms/step - acc: 0.9448 - loss: 0.4658
[0.42359066009521484, 0.9580419659614563]
```

Indeed, the accuracy of the model improved from approximately 0.9161 (91.61%) before normalization to approximately 0.9650 (96.50%) after normalization

Use Image Classification on the hand written digits data set (mnist)

- 1. Plot the loss and accuracy for both training data and validation data using the history object in the source code.
- 2. Plot one of the images in the test data, and then do inferencing to check what is the prediction of the model on that single image.

```
from keras import Sequential
from keras.datasets import mnist
import numpy as np
from keras.layers import Dense
from keras.utils import to categorical
import matplotlib.pyplot as plt
(train images, train labels), (test images, test labels) = mnist.load data()
print(train images.shape[1:])
#process the data
#1. convert each image of shape 28*28 to 784 dimensional which will be fed to the network as a single feature
dimData = np.prod(train_images.shape[1:])
print(dimData)
train data = train images.reshape(train images.shape[0],dimData)
test_data = test_images.reshape(test_images.shape[0],dimData)
#convert data to float and scale values between 0 and 1
train data = train data.astype('float')
test_data = test_data.astype('float')
#scale data
train_data /=255.0
test_data /=255.0
#change the labels frominteger to one-hot encoding. to_categorical is doing the same thing as LabelEncoder()
train_labels_one_hot = to_categorical(train_labels)
test labels one hot = to_categorical(test_labels)
```

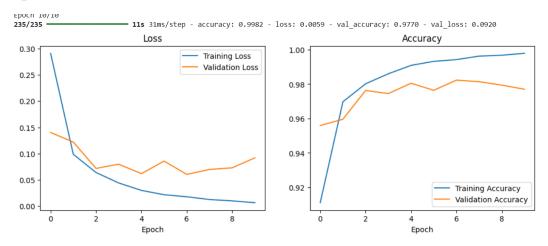
```
#creating network
model = Sequential()
model.add(Dense(512, activation='relu', input_shape=(dimData,)))
model.add(Dense(512, activation='relu'))
model.add(Dense(10, activation='softmax'))

model.compile(optimizer='rmspror'_loss-'categorical_crossentropy', metrics=['accuracy'])
history = model.fit(train_data, loading... els_one_hot, batch_size=256, epochs=10, validation_data=(test_data, test_labels_one_hot))

# Extract training history
training_loss = history.history['loss']
training_accuracy = history.history['accuracy']
validation_loss = history.history['val_loss']
validation_accuracy = history.history['val_accuracy']
```

```
# Plot loss
plt.figure(figsize=(12, 4))
plt.subplot(1, 2, 1)
plt.plot(training loss, label='Training Loss')
plt.plot(validation_loss, label='Validation Loss')
plt.title('Loss')
plt.xlabel('Epoch')
plt.legend()
# Plot accuracy
plt.subplot(1, 2, 2)
plt.plot(training_accuracy, label='Training Accuracy')
plt.plot(validation_accuracy, label='Validation Accuracy')
plt.title('Accuracy')
plt.xlabel('Epoch')
plt.legend()
plt.show()
```

Output:



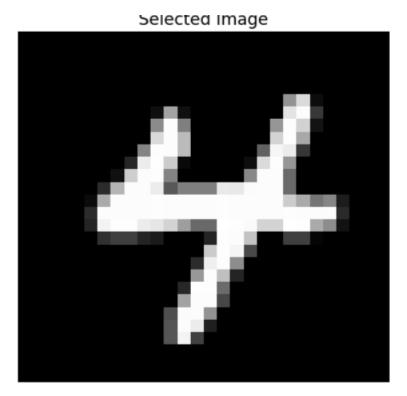
2. Plot one of the images in the test data, and then do inferencing to check what is the prediction of the model on that single image.

```
# plot the selected image
plt.figure()
plt.imshow(image, cmap='gray')
plt.axis('off')
plt.title('Selected Image')

# do inferencing to check the model prediction on the selected image
prediction = model.predict(image.reshape(1, 784))
prediction = np.argmax(prediction)

# print the predicted label
print('Predicted label:', prediction)
```

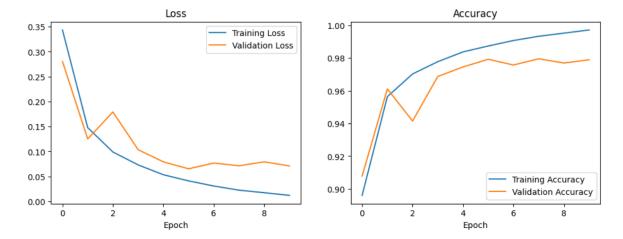
Output:



3. We had used 2 hidden layers and Relu activation. Try to change the number of hidden layer and the activation to tanh or sigmoid and see what happens.

```
[ ] # 3. We had used 2 hidden layers and Relu activation. Try to change the number of hidden layer and the
        #activation to tanh or sigmoid and see what happens.
[ ] from keras import Sequential
    from keras.datasets import mnist
    import numpy as np
    from keras.layers import Dense
from keras.utils import to_categorical
    import matplotlib.pyplot as plt
    (train_images,train_labels),(test_images, test_labels) = mnist.load_data()
    print(train images.shape[1:])
    #process the data
    #1. convert each image of shape 28*28 to 784 dimensional which will be fed to the network as a single feature
    dimData = np.prod(train_images.shape[1:])
    print(dimData)
    train_data = train_images.reshape(train_images.shape[0],dimData)
    test_data = test_images.reshape(test_images.shape[0],dimData)
    #convert data to float and scale values between 0 and 1
    train_data = train_data.astype('float')
    test_data = test_data.astype('float')
    #scale data
    train_data /=255.0
    test data /=255.0
    #change the labels frominteger to one-hot encoding. to_categorical is doing the same thing as LabelEncoder()
    train labels one hot = to categorical(train labels)
    test_labels_one_hot = to_categorical(test_labels)
  #creating network
   model = Sequential()
   model.add(Dense(512, activation='tanh', input_shape=(dimData,)))
   model.add(Dense(256, activation='tanh'))
   model.add(Dense(128, activation='tanh'))
   model.add(Dense(10, activation='softmax'))
   model.compile(optimizer='rmsprop', loss='categorical_crossentropy', metrics=['accuracy'])
   history = model.fit(train_data, train_labels_one_hot, batch_size=256, epochs=10, verbose=1,
                      validation_data=(test_data, test_labels_one_hot))
   # Extract training history
   training_loss = history.history['loss']
   training_accuracy = history.history['accuracy']
   validation loss = history.history['val loss']
   validation_accuracy = history.history['val_accuracy']
     # Plot loss
      plt.figure(figsize=(12, 4))
      plt.subplot(1, 2, 1)
      plt.plot(training_loss, label='Training Loss')
      plt.plot(validation_loss, label='Validation Loss')
      plt.title('Loss')
      plt.xlabel('Epoch')
      plt.legend()
      # Plot accuracy
      plt.subplot(1, 2, 2)
plt.plot(training_accuracy, label='Training Accuracy')
      plt.plot(validation_accuracy, label='Validation Accuracy')
      plt.title('Accuracy'
      plt.xlabel('Epoch')
      plt.legend()
      plt.show()
```

Output:



4. Run the same code without scaling the images and check the performance?

```
import matplotlib.pyplot as plt
from keras import Sequential
from keras import Sequential
from keras datasets import mumpy as np
from keras.layers import Dense
from keras.layers import to categorical

(train_images, train_labels), (test_images, test_labels) = mnist.load_data()

print(train_images.shape[1:])
# Process the data
# 1. Convert each image of shape 28*28 to 784 dimensional which will be fed to the network as a single feature dimData = np.prod(train_images.shape[1:])
print(dimData)
train_data = train_images.reshape(train_images.shape[0], dimData)

# Convert data to float (no scaling)
train_data = train_data.astype('float')
test_data = test_data.astype('float')

# Change the labels from integer to one-hot encoding. to_categorical is doing the same thing as LabelEncoder()
train_labels_one_hot = to_categorical(train_labels)
test_labels_one_hot = to_categorical(test_labels)
```

```
# Creating network
model = Sequential()
model.add(Dense(512, activation='tanh', input_shape=(dimData,)))
model.add(Dense(256, activation='tanh'))
model.add(Dense(128, activation='tanh'))
model.add(Dense(10, activation='softmax'))
model.compile(optimizer='rmsprop', loss='categorical_crossentropy', metrics=['accuracy'])
history = model.fit(train_data__train_labels_one_hot, batch_size=256, epochs=10, verbose=1,
                    validation_data: Any data, test_labels_one_hot))
# Extract training history
training_loss = history.history['loss']
training_accuracy = history.history['accuracy']
validation_loss = history.history['val_loss']
validation_accuracy = history.history['val_accuracy']
# Plot loss
plt.figure(figsize=(12, 4))
plt.subplot(1, 2, 1)
plt.plot(training_loss, label='Training Loss')
plt.plot(validation_loss, label='Validation Loss')
plt.title('Loss')
plt.xlabel('Epoch')
plt.legend()
# Plot accuracy
plt.subplot(1, 2, 2)
plt.plot(training_accuracy, label='Training Accuracy')
plt.plot(validation_accuracy, label='Validation Accuracy')
plt.title('Accuracy')
plt.xlabel('Epoch')
plt.legend()
plt.show()
```

Output:

