Spring 2024: CS5720 Neural Networks & Deep Learning - ICP-6 Bhanu Chandrika Lakkimsetti (700747439)

ICP_Basics in Keras

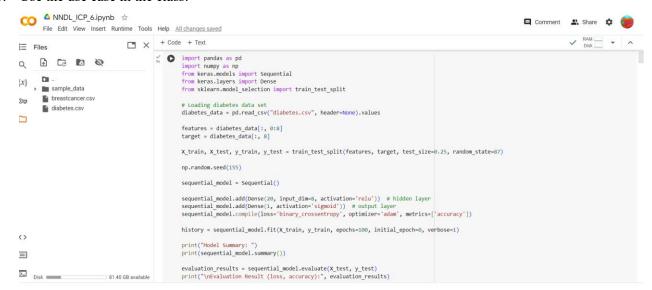
GitHub Link: https://github.com/bhanuchandrika99/NNDL ICP 6

Use Case Description: Predicting the diabetes disease

Programming elements: Keras Basics

In class programming:

1. Use the use case in the class:



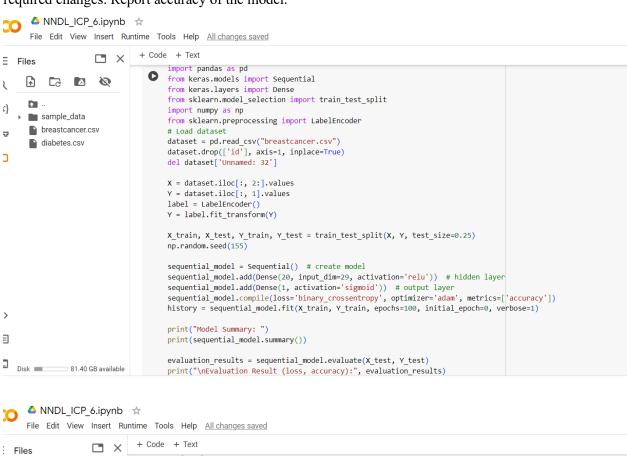
```
0
 Epoch 1/100
\Box
 Epoch 2/100
        18/18 [======
 Epoch 3/100
 Epoch 4/100
 18/18 [=============] - Øs 2ms/step - loss: 7.7556 - accuracy: 0.6007
 Epoch 5/100
        18/18 [=====
 Epoch 6/100
 18/18 [======
         Epoch 7/100
 Epoch 8/100
 18/18 [==============] - 0s 2ms/step - loss: 3.0615 - accuracy: 0.5417
 Epoch 9/100
 Enoch 10/100
 Epoch 11/100
 Epoch 12/100
 18/18 [============] - Os 2ms/step - loss: 0.9967 - accuracy: 0.6389
 Epoch 13/100
 18/18 [======
        ============ ] - Os 2ms/step - loss: 0.8826 - accuracy: 0.6615
 Epoch 14/100
 18/18 [============] - 0s 2ms/step - loss: 0.8023 - accuracy: 0.6684
```

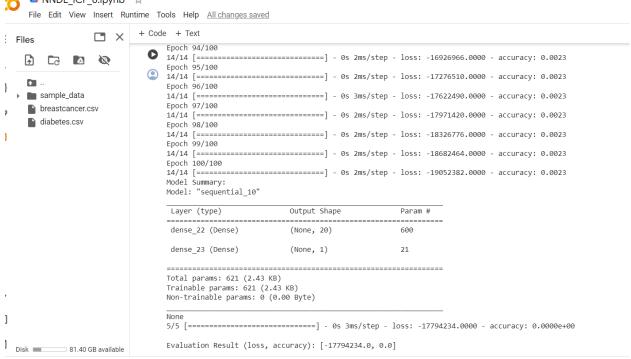
```
+ Code + Text
    P Epoch 95/100
    18/18 [=============] - 0s 3ms/step - loss: 0.6055 - accuracy: 0.7066
 Epoch 96/100
    Epoch 97/100
    18/18 [==============] - 0s 2ms/step - loss: 0.5554 - accuracy: 0.7188
    Epoch 98/100
    Epoch 99/100
    18/18 [==============] - 0s 2ms/step - loss: 0.5734 - accuracy: 0.7118
    Epoch 100/100
    18/18 [============] - 0s 2ms/step - loss: 0.6140 - accuracy: 0.6944
    Model Summary:
    Model: "sequential 3"
    Layer (type)
                      Output Shape
                                       Param #
    ______
    dense_10 (Dense)
                       (None, 20)
    dense_11 (Dense)
                      (None, 1)
    Total params: 201 (804.00 Byte)
    Trainable params: 201 (804.00 Byte)
    Non-trainable params: 0 (0.00 Byte)
    6/6 [============] - 0s 3ms/step - loss: 0.6438 - accuracy: 0.6823
    Evaluation Result (loss, accuracy): [0.6437555551528931, 0.6822916865348816]
```

a. Add more Dense layers to the existing code and check how the accuracy changes.

```
△ NNDL_ICP_6.ipynb ☆
File Edit View Insert Runtime Tools Help All changes saved
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ì
   C
            Ø
                   [3] import pandas as pd
                Show hidden files m keras. models import Sequential m keras.layers import Dense
sample_data
                         from sklearn.model_selection import train_test_split
breastcancer.csv
                         import numpy as np
diabetes.csv
                         # Load dataset
                         dataset = pd.read_csv("diabetes.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)
                         sequential_model = Sequential() # create model
                         sequential_model.add(Dense(20, input_dim=8, activation='relu'))
                         sequential_model.add(Dense(10, activation='relu')) # Additional hidden layer
                         sequential_model.add(Dense(5, activation='relu'))
                                                                # Additional hidden layer
                         sequential_model.add(Dense(1, activation='sigmoid')) # output layer
                         sequential_model.compile(loss='binary_crossentropy', optimizer='adam', metrics=[|'accuracy'])
                         history = sequential_model.fit(X_train, Y_train, epochs=100,
                                                     initial_epoch=0, verbose=1)
                         print("Model Summary: ")
                         print(sequential model.summary())
                         evaluation results = sequential model.evaluate(X test, v test)
                         print("\nEvaluation Result (loss, accuracy):", evaluation_results)
81.40 GB available
  + Code + Text
        TO/ TO [ --
                               Epoch 97/100
        18/18 [============= ] - 0s 2ms/step - loss: 0.5674 - accuracy: 0.7101
    ➡ Epoch 98/100
        18/18 [============= ] - 0s 2ms/step - loss: 0.5670 - accuracy: 0.7118
        Epoch 99/100
        18/18 [=============] - 0s 2ms/step - loss: 0.5666 - accuracy: 0.7153
        Epoch 100/100
        Model Summary:
        Model: "sequential_1"
         Layer (type)
                                  Output Shape
                                                          Param #
        ______
         dense 2 (Dense)
                                   (None, 20)
                                                          180
         dense 3 (Dense)
                                   (None, 10)
                                                          210
         dense 4 (Dense)
                                   (None, 5)
                                                          55
         dense_5 (Dense)
                                   (None, 1)
                                                          6
        _____
        Total params: 451 (1.76 KB)
        Trainable params: 451 (1.76 KB)
        Non-trainable params: 0 (0.00 Byte)
        None
        Evaluation Result (loss, accuracy): [0.6229787468910217, 0.6458333134651184]
```

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

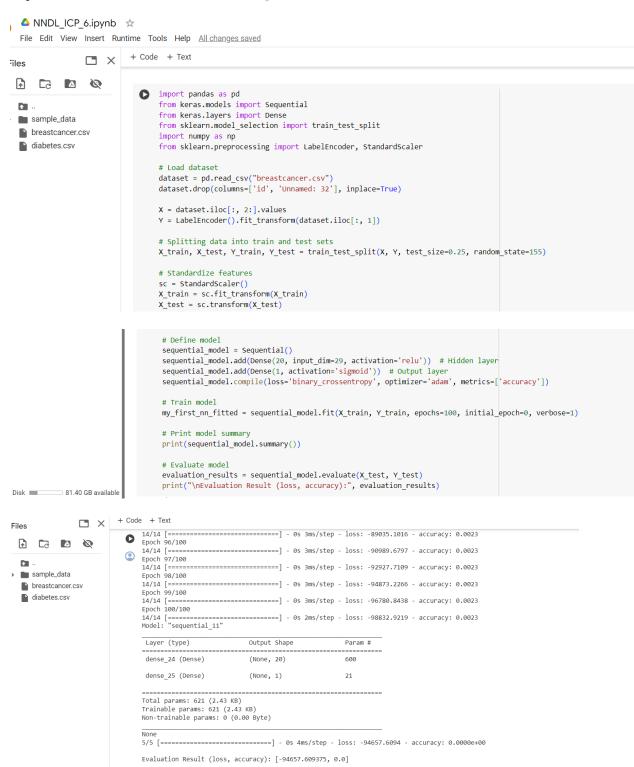




3. Normalize the data before feeding the data to the model and check how the normalization change your accuracy (code given below).

from sklearn.preprocessing

import StandardScaler sc = StandardScaler()



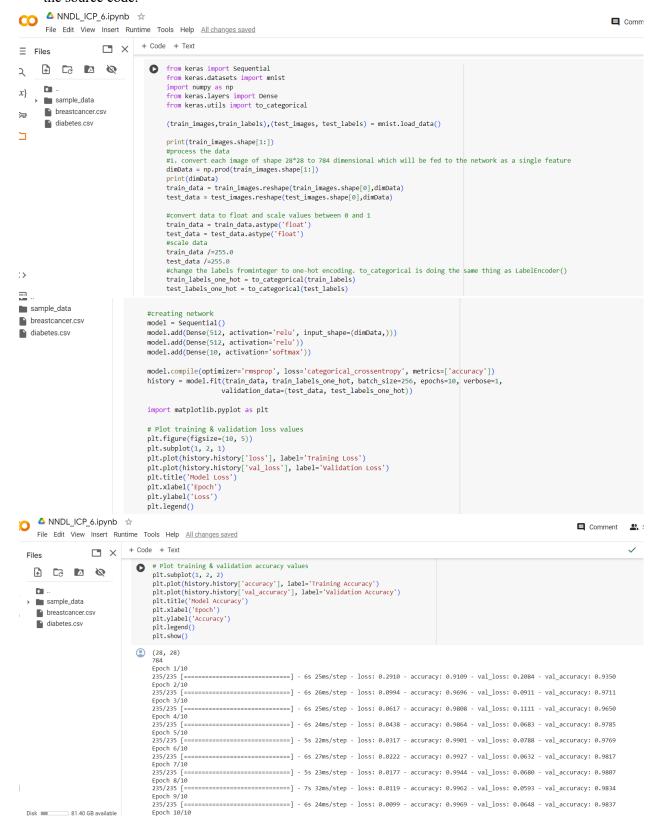
Note: Breast Cancer dataset is designated to predict if a patient has Malignant (M) or Benign = B cancer

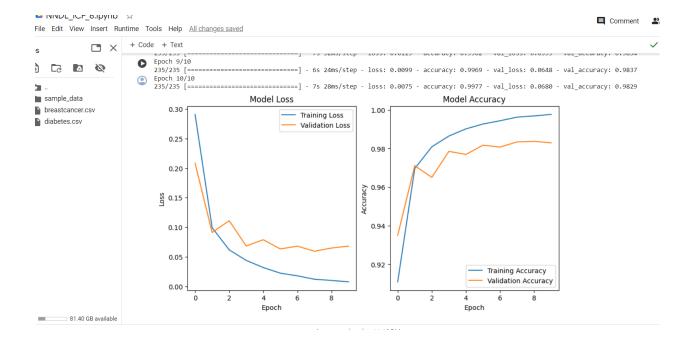
In class programming:

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

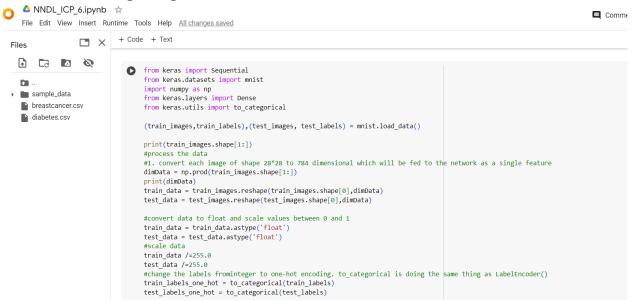
```
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                   □ × + Code + Text
≔ Files
    from keras import Sequential
                                 from keras datasets import mnist
     CII ...
                                 import numpy as np
   sample_data
                                 from keras.utils import to_categorical
    breastcancer.csv
      diabetes.csv
                                 (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.astvpe('float')
                                 test_data = test_data.astype('float')
                                 #scale data
                                 train_data /=255.0
                                  test_data /=255.0
                                 #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)
\equiv
sample_data
                            #creating network
breastcancer.csv
                            model = Sequential()
diabetes.csv
                           model.add(Dense(512, activation='relu', input_shape=(dimData,)))
model.add(Dense(512, activation='relu'))
                            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))
 breastcancer.csv
                        Downloading data from <a href="https://storage.googleapis.com/tensorflow/tf-keras-datasets/mnist.npz">https://storage.googleapis.com/tensorflow/tf-keras-datasets/mnist.npz</a>
 diabetes.csv
                            11490434/11490434 [
                            (28, 28)
                            Epoch 1/10
                            235/235 [=
Epoch 2/10
                                                    235/235 [==:
                                        ==========] - 8s 33ms/step - loss: 0.0627 - accuracy: 0.9806 - val_loss: 0.0905 - val_accuracy: 0.9709
                            Epoch 4/10
                            235/235 [===
Epoch 5/10
                                           =========================== - 6s 25ms/step - loss: 0.0434 - accuracy: 0.9864 - val_loss: 0.0602 - val_accuracy: 0.9812
                            =========] - 6s 24ms/step - loss: 0.0221 - accuracy: 0.9930 - val_loss: 0.0661 - val_accuracy: 0.9811
                            Epoch 7/10
                            235/235 [===
                                             =========] - 6s 27ms/step - loss: 0.0162 - accuracy: 0.9948 - val_loss: 0.0634 - val_accuracy: 0.9818
                            235/235 [==========] - 5s 23ms/step - loss: 0.0125 - accuracy: 0.9960 - val loss: 0.0748 - val accuracy: 0.9801
                            Fnoch 9/10
                                        235/235 [====
```

1. Plot the loss and accuracy for both training data and validation data using the history object in the source code.

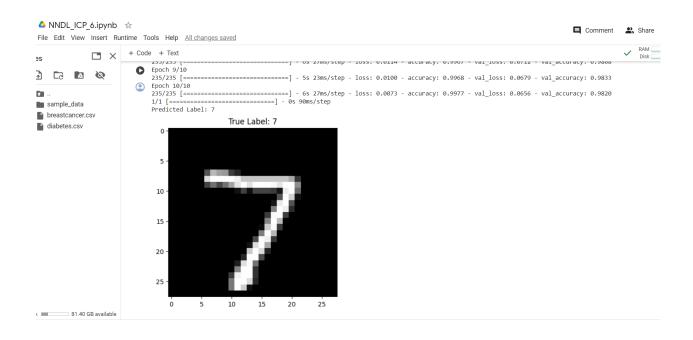




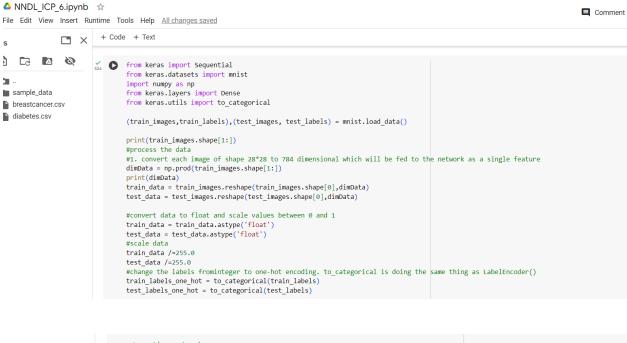
2. Plot one of the images in the test data, and then do inferencing to check what is the prediction of the model on that single image.







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.



235/235 [================================] - 5s 21ms/step - loss: 0.0125 - accuracy: 0.9969 - val_loss: 0.0923 - val_accuracy: 0.9762

```
breastcancer.csv
diabetes.csv
              (28, 28)
                       ========== ] - 5s 21ms/step - loss: 0.3397 - accuracy: 0.8967 - val loss: 0.1980 - val accuracy: 0.9384
              235/235 [==
              Epoch 2/10
                       Epoch 3/10
              235/235 [==
                       Epoch 4/10
                     Epoch 5/10
              235/235 [============] - 5s 20ms/step - loss: 0.0540 - accuracy: 0.9827 - val_loss: 0.0936 - val_accuracy: 0.9702 
Epoch 6/10
                       Epoch 7/10
              235/235 [============] - 5s 23ms/step - loss: 0.0309 - accuracy: 0.9911 - val_loss: 0.0907 - val_accuracy: 0.9707 Epoch 8/10
```

sample_data

Epoch 10/10

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

