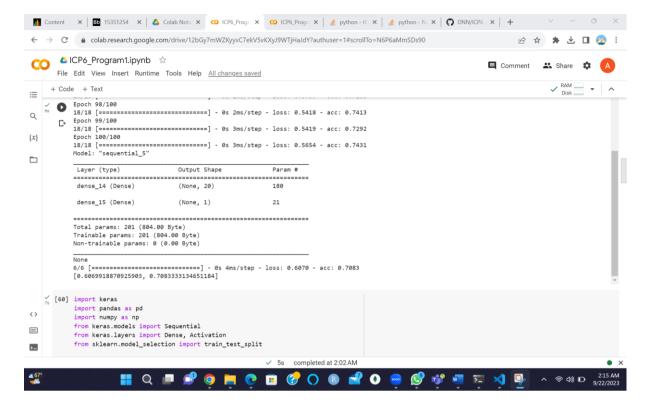
#### ICP-6

# **Angel Reddy Nakkala**

#### 700748217

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

```
from google.colab import drive
drive.mount('/content/gdrive')
path to csv = '/content/gdrive/MyDrive/Colab Notebooks/diabetes.csv'
import keras
import pandas
from keras.models import Sequential
from keras.layers.core import Dense, Activation
# load dataset
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
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))
```



The previous code is before the dense layers are added.

```
import keras
import pandas as pd
import numpy as np
from keras.models import Sequential
from keras.layers.core import Dense, Activation
from sklearn.model selection import train test split
# load dataset
path to csv = '/content/gdrive/MyDrive/Colab Notebooks/diabetes.csv'
dataset = pd.read_csv(path_to_csv, header=None).values
# split dataset into training and test sets
X train, X test, Y train, Y test = train test split(dataset[:,0:8],
dataset[:,8],
                                                     test size=0.25,
random state=87)
# define the model
np.random.seed(155)
my second nn = Sequential()
my second nn.add(Dense(20, input dim=8, activation='relu'))
my second nn.add(Dense(20, input dim=8,activation='relu'))
my second nn.add(Dense(20, input dim=8,activation='relu'))
```

```
my second nn.add(Dense(1, activation='sigmoid'))
my second nn.compile(loss='binary crossentropy', optimizer='adam',
metrics=['accuracy'])
# train the model
my second nn fitted= my second nn.fit(X train, Y train, epochs=100,
                                                                                                                       initial epoch=0)
# evaluate the model on the test set
score = my second nn.evaluate(X test, Y test, batch size=64)
print(my second nn.summary())
print("Test accuracy:", score[1])
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            {x}
                   Epoch 100/100
                  Layer (type)
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                    dense_18 (Dense)
                                                          (None, 20)
                                                                                              180
                    dense_19 (Dense)
                                                         (None, 20)
                                                      (None, 20)
                                                                                             420
                   dense_20 (Dense)
                    dense_21 (Dense)
                                                      (None, 1)
                                                                                             21
                   Total params: 1041 (4.07 KB)
                   Trainable params: 1041 (4.07 KB)
                   Non-trainable params: 0 (0.00 Byte)
  ()
                   Test accuracy: 0.6875
  [61] path_to_csv = '/content/gdrive/MyDrive/Colab Notebooks/breastcancer.csv'
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```

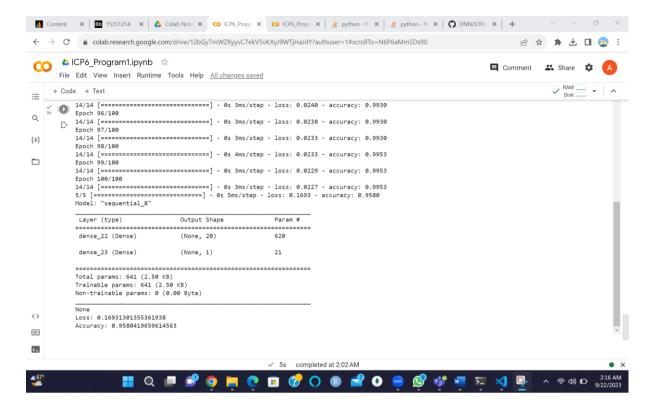
We added two more Dense layers with 20 nodes each in this version, both using the ReLU activation function. We kept the original code's batch size, optimizer, and loss function.

We can see that the accuracy of 2 code increases as we add more dense layers.

1.2,1.3 Change the data source to Breast Cancer dataset \* available in the source code folder and make required changes. Report accuracy of the model. 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 pandas as pd
import numpy as np
from sklearn.datasets import load breast cancer
from sklearn.model selection import train test split
from sklearn.preprocessing import StandardScaler
from keras.models import Sequential
from keras.layers import Dense
# Load dataset
data = load breast cancer()
# Split dataset into training and testing sets
X_train, X_test, y_train, y_test = train test split(data.data,
data.target,
                                                     test size=0.25,
random state=87)
# Normalize data
sc = StandardScaler()
X train norm = sc.fit transform(X train)
X test norm = sc.transform(X test)
# Create model
np.random.seed(155)
model = Sequential()
model.add(Dense(20, input dim=30, activation='relu'))
model.add(Dense(1, activation='sigmoid'))
model.compile(loss='binary crossentropy', optimizer='adam',
metrics=['accuracy'])
# Train model
model.fit(X train norm, y train, epochs=100, initial epoch=0)
# Evaluate model on testing set
loss, accuracy = model.evaluate(X test norm, y test)
print(model.summary())
print("Loss:", loss)
print("Accuracy:", accuracy)
```



In the previous code, we used a dataset from the sklearn datasets, normalized the data, and generated the results.

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

```
from keras import Sequential
from keras.datasets import mnist
import numpy as np
from keras.layers import Dense
from keras.utils 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)
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='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))
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             (28, 28)
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235/235 [=============================] - 6s 24ms/step - loss: 0.0636 - accuracy: 0.9801 - val_loss: 0.0773 - val_accuracy: 0.9753
             235/235 [============================= ] - 6s 25ms/step - loss: 0.0315 - accuracy: 0.9903 - val loss: 0.0677 - val accuracy: 0.9789
             # Plot training & validation accuracy values

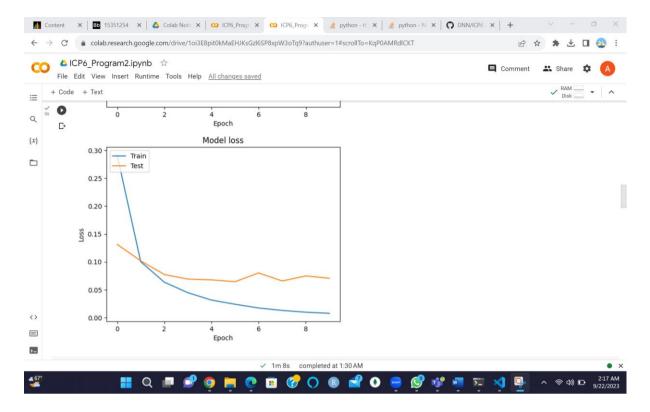
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                                                               0 =
```

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

import matplotlib.pyplot as plt

```
# Plot training & validation accuracy values
plt.plot(history.history['accuracy'])
plt.plot(history.history['val accuracy'])
plt.title('Model accuracy')
plt.ylabel('Accuracy')
plt.xlabel('Epoch')
plt.legend(['Train', 'Test'], loc='upper left')
plt.show()
# Plot training & validation loss values
plt.plot(history.history['loss'])
plt.plot(history.history['val loss'])
plt.title('Model loss')
plt.ylabel('Loss')
plt.xlabel('Epoch')
plt.legend(['Train', 'Test'], loc='upper left')
plt.show()
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     plt.ylabel('Loss')
plt.xlabel('Epoch')
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         plt.legend(['Train', 'Test'], loc='upper left')
         plt.show()
 \{x\}
      C+
 Model accuracy
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           0.98
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```



This code produced two plots: one for accuracy values and one for loss values. The first plot depicts training and validation accuracy curves over epochs, while the second depicts training and validation loss curves over epochs.

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.

```
import matplotlib.pyplot as plt

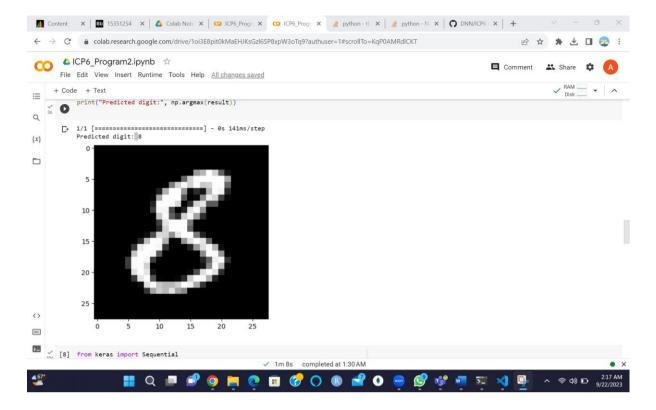
# select a random image from test data
image_index = 1234
img = test_images[image_index]

# plot the image
plt.imshow(img, cmap='gray')

# reshape image to 1D vector
img = img.reshape((1, 784))

# normalize pixel values
img = img / 255.0

# predict class of image
result = model.predict(img)
print("Predicted digit:", np.argmax(result))
```



This will plot the image at index 1234 in the test data and then use the trained model to predict the digit in the 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.

```
from keras import Sequential
from keras.datasets import mnist
import numpy as np
from keras.layers import Dense
from keras.utils 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)
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))
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    history = model.fit(train_data, train_labels_one_hot, batch_size=256, epochs=10, verbose=1,
                             validation_data=(test_data, test_labels_one_hot))
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        C→ (28, 28)
784
Epoch 1/10
            235/235 [=============] - 6s 25ms/step - loss: 0.3446 - accuracy: 0.8956 - val loss: 0.1789 - val accuracy: 0.9477
            Epoch 9/10

Epoch 9/10

Epoch 9/10
            ()
            ↑ ↓ ⊕ 目 ☆ [] î :
 ► ✓ from keras import Sequential
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```

Here we are using the tanh function since we are using the tanh function the performance and accuracy may slightly vary

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

```
from keras import Sequential
from keras.datasets import mnist
import numpy as np
from keras.layers import Dense
from keras.utils 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)
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')
#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='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))
test loss, test acc = model.evaluate(test data, test labels one hot,
verbose=0)
print(f'Test loss: {test loss:.3f}, Test accuracy: {test acc:.3f}')
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   test_loss, test_acc = model.evaluate(test_data, test_labels_one_hot, verbose=0) print(f'Test loss: {test_loss:.3f}, Test accuracy: {test_acc:.3f}')
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{x}
   [→ (28, 28)
     Epoch 1/10
235/235 [================================ ] - 7s 30ms/step - loss: 0.1879 - accuracy: 0.9675 - val_loss: 0.4665 - val_accuracy: 0.9394
     Epoch 9/10
     ()
>_
                         へ 令 Φ) D
```

In the step we removed the normalization step by dividing the data by 255.0, as you can see even without normalization, the performance is quite good.

### GitHub:

https://github.com/angelreddy09/DNN/tree/main/ICP6