# CS 5720 Neural Network Deep Learning ICP-7

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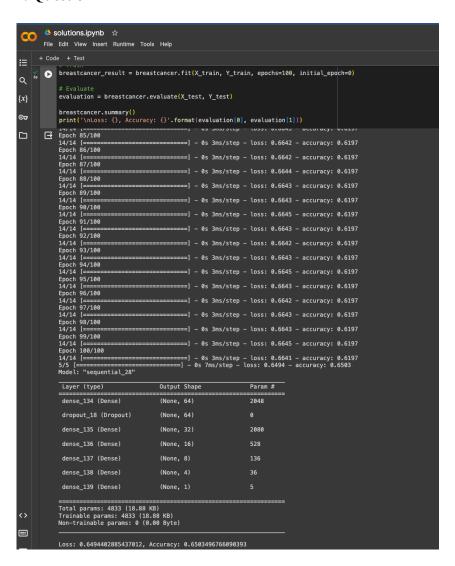
#### **GitHub Repository:**

https://github.com/mxb40210/700754021-NeuralNetworkDeepLearning

### **Assignment 7:**

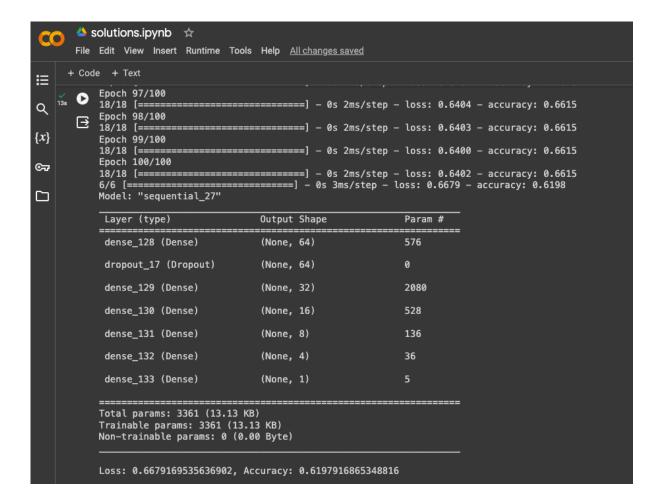
https://github.com/mxb40210/700754021-NeuralNetworkDeepLearning/tree/main/assignments/assignment7

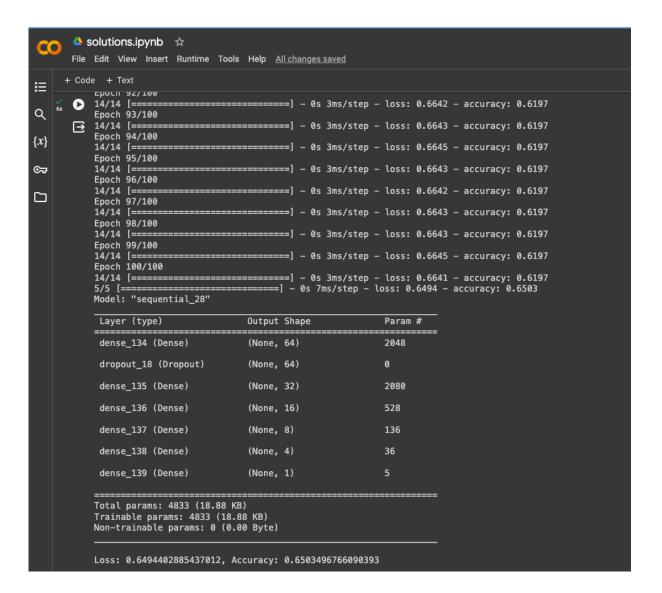
#### 1. Question 1



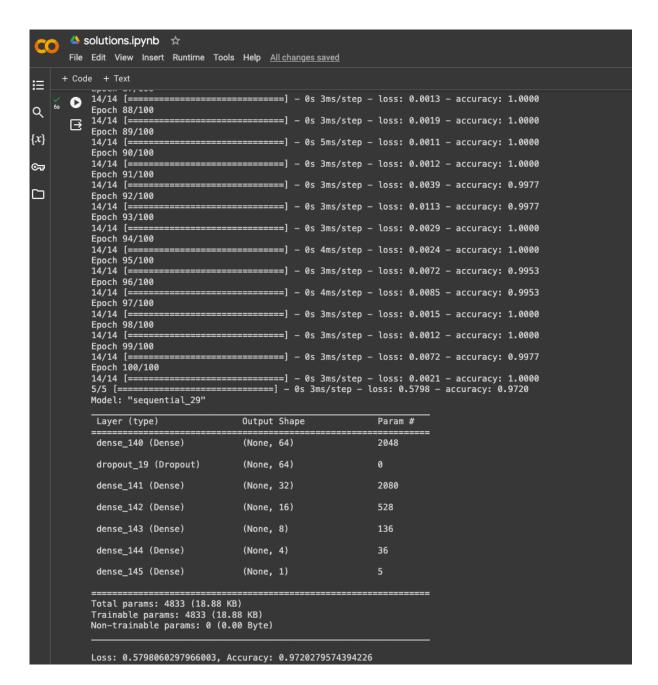
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Q / 13s D """
                1.1 Use the use case in the class:
a. Add more Dense layers to the existing code and check how the accuracy changes.
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                diabetes_dataset = pd.read_csv(diabetes_csv_path, header=None).values
# Split the dataset X and Y
X = diabetes_dataset[:, 0:8]
                 Y = diabetes_dataset[:, 8]
                X_train, X_test, Y_train, Y_test = train_test_split(X, Y, test_size=0.25, random_state=87)
                np.random.seed(155)
                 # Create model
                diabetes_model = Sequential()
                # Add layers
diabetes_model.add(Dense(64, input_dim=8, activation='relu')) # Input layer
diabetes_model.add(Dropout(0.2)) # Dropout layer to prevent overfitting
diabetes_model.add(Dense(32, activation='relu')) # Hidden layer
diabetes_model.add(Dense(16, activation='relu')) # Hidden layer
diabetes_model.add(Dense(8, activation='relu')) # Hidden layer
diabetes_model.add(Dense(4, activation='relu')) # Hidden layer
diabetes_model.add(Dense(1, activation='relu')) # Output layer
                 sgd = SGD(learning_rate=0.01, momentum=0.9)
                 # Compile
                diabetes_model.compile(loss='binary_crossentropy', optimizer=sgd, metrics=['accuracy'])
                 diabetes_result = diabetes_model.fit(X_train, Y_train, epochs=100, initial_epoch=0)
                 # Evaluate
                 evaluation = diabetes_model.evaluate(X_test, Y_test)
                 diabetes_model.summary()
                 print('\nLoss: {}, Accuracy: {}'.format(evaluation[0], evaluation[1]))
```

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Q 6s D """
               1.2 Change the data source to Breast Cancer dataset * available in the source code folder and make required changes. Report accuracy of the model.
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                # Read breastcancer dataset
                breastcancer_dataset = pd.read_csv(breastcancer_csv_path)
breastcancer_dataset = breastcancer_dataset.dropna(axis=1)
                # Map target labels to 'M' and 'B'
label_encoder = LabelEncoder()
                breastcancer_dataset['diagnosis'] = label_encoder.fit_transform(breastcancer_dataset['diagnosis'])
                # # Split the dataset X and Y
               X = breastcancer_dataset.drop('diagnosis', axis=1)
Y = breastcancer_dataset['diagnosis']
                X_train, X_test, Y_train, Y_test = train_test_split(X, Y, test_size=0.25, random_state=87)
                np.random.seed(155)
                # Create model
                breastcancer = Sequential()
               # Add layers
breastcancer.add(Dense(64, input_dim=X.shape[1], activation='relu')) # Input layer
breastcancer.add(Dropout(0.2)) # Dropout layer to prevent overfitting
breastcancer.add(Dense(32, activation='relu')) # Hidden layer
breastcancer.add(Dense(16, activation='relu')) # Hidden layer
breastcancer.add(Dense(8, activation='relu')) # Hidden layer
breastcancer.add(Dense(4, activation='relu')) # Hidden layer
breastcancer.add(Dense(1, activation='relu')) # Output layer
                # Optimizer
                sgd = SGD(learning_rate=0.01, momentum=0.9)
                breastcancer.compile(loss='binary_crossentropy', optimizer=sgd, metrics=['accuracy'])
                breastcancer_result = breastcancer.fit(X_train, Y_train, epochs=100, initial_epoch=0)
                evaluation = breastcancer.evaluate(X_test, Y_test)
                breastcancer.summary()
print('\nLoss: {}, Accuracy: {}'.format(evaluation[0], evaluation[1]))
```

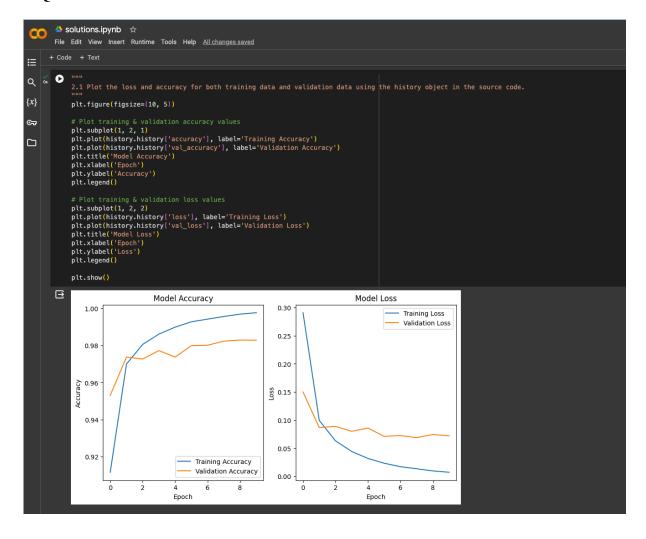




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                  import StandardScale
                  sc = StandardScaler()
# Read breastcancer dataset
                 breastcancer_dataset = pd.read_csv(breastcancer_csv_path)
breastcancer_dataset = breastcancer_dataset.dropna(axis=1)
                 label_encoder = LabelEncoder()
breastcancer_dataset['diagnosis'] = label_encoder.fit_transform(breastcancer_dataset['diagnosis'])
                  sc = StandardScaler()
X_normalized = sc.fit_transform(breastcancer_dataset.drop('diagnosis', axis=1))
                  X = X_normalized
Y = breastcancer_dataset['diagnosis']
                  X_train, X_test, Y_train, Y_test = train_test_split(X, Y, test_size=0.25, random_state=87)
                  # Set some seed
np.random.seed(155)
                  # Create model
breastcancer = Sequential()
                 # Add layers
breastcancer.add(Dense(64, input_dim=X.shape[1], activation='relu')) # Input layer
breastcancer.add(Dense(62)) # Dropout layer to prevent overfitting
breastcancer.add(Dense(32, activation='relu')) # Hidden layer
breastcancer.add(Dense(16, activation='relu')) # Hidden layer
breastcancer.add(Dense(8, activation='relu')) # Hidden layer
breastcancer.add(Dense(4, activation='relu')) # Hidden layer
breastcancer.add(Dense(1, activation='rslu')) # Output layer
                  # Optimize
                  sgd = SGD(learning_rate=0.01, momentum=0.9)
                  # Compile
breastcancer.compile(loss='binary_crossentropy', optimizer=sgd, metrics=['accuracy'])
                  breastcancer_result = breastcancer.fit(X_train, Y_train, epochs=100, initial_epoch=0)
                  # Evaluate
evaluation = breastcancer.evaluate(X_test, Y_test)
                  breastcancer.summary()
print('\nLoss: {}, Accuracy: {}'.format(evaluation[0], evaluation[1]))
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## 2. Question 2



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   Epoch 5/10
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Epoch 6/10
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Epoch 7/10
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           {x}
    Epoch 8/10
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Epoch 9/10
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          :========] - 9s 40ms/step - loss: 0.0138 - accuracy: 0.9959 - val_loss: 0.0782 - val_accuracy: 0.9762
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Epoch 3/10
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Epoch 4/10
           :=========] - 7s 31ms/step - loss: 0.2993 - accuracy: 0.9100 - val_loss: 0.3031 - val_accuracy: 0.9091
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    235/235 [==
Epoch 5/10
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Epoch 6/10
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Epoch 7/10
            =========] - 8s 33ms/step - loss: 0.2067 - accuracy: 0.9375 - val_loss: 0.1904 - val_accuracy: 0.9412
           235/235
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{x}
                   # Load the MNIST datase
                  (train_images, train_labels), (test_images, test_labels) = mnist.load_data()
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                  # Process the data
dimData = np.prod(train_images.shape[1:])
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                  train_data = train_images.reshape(train_images.shape[0], dimData).astype('float32')
test_data = test_images.reshape(test_images.shape[0], dimData).astype('float32')
train_labels.one_hot = to_categorical(train_labels)
test_labels_one_hot = to_categorical(test_labels)
                  for num_layers in num_hidden_layers:
                             validation_data=(test_data, test_labe

Factor in property in the process of the 
                                              ==========] - 3s 13ms/step - loss: 0.2493 - accuracy: 0.9246 - val_loss: 0.2308 - val_accuracy: 0.9334
                                                            ========] - 3s 13ms/step - loss: 0.1368 - accuracy: 0.9582 - val_loss: 0.1449 - val_accuracy: 0.9576
                                                                :=======] - 5s 19ms/step - loss: 0.1307 - accuracy: 0.9599 - val_loss: 0.1425 - val_accuracy: 0.9562
                                                    ========] - 4s 19ms/step - loss: 0.1856 - accuracy: 0.9449 - val_loss: 0.1724 - val_accuracy: 0.9488
                                                                 ========] - 3s 13ms/step - loss: 0.1639 - accuracy: 0.9511 - val_loss: 0.1556 - val_accuracy: 0.9546
                                                              =======] - 3s 13ms/step - loss: 0.1339 - accuracy: 0.9597 - val_loss: 0.1364 - val_accuracy: 0.9589
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