

Configuration Manual

MSc Research Project
Programme Name

Anthony Effiok
x19108788

School of Computing
National College of Ireland

Supervisor: Dr. Hicham Rifai

National College of Ireland
MSc Project Submission Sheet



School of Computing

Student Name: Anthony Eyo Effiok

Student ID: X19108788

Programme: Msc Data Analytics **Year:** 2020

Module: Research Project

Lecturer: Dr. Hicham Rifai

Submission Due Date: 17TH August 2020

Project Title: NiohSign: A Siamese neural network approach for Signature Authentication

Word Count: 685 **Page Count:** 11

I hereby certify that the information contained in this (my submission) is information pertaining to research I conducted for this project. All information other than my own contribution will be fully referenced and listed in the relevant bibliography section at the rear of the project.

ALL internet material must be referenced in the bibliography section. Students are required to use the Referencing Standard specified in the report template. To use other author's written or electronic work is illegal (plagiarism) and may result in disciplinary action.

Signature: Anthony Eyo Effiok

Date: 17TH August 2020

PLEASE READ THE FOLLOWING INSTRUCTIONS AND CHECKLIST

Attach a completed copy of this sheet to each project (including multiple copies)	<input type="checkbox"/>
Attach a Moodle submission receipt of the online project submission, to each project (including multiple copies).	<input type="checkbox"/>
You must ensure that you retain a HARD COPY of the project, both for your own reference and in case a project is lost or mislaid. It is not sufficient to keep a copy on computer.	<input type="checkbox"/>

Assignments that are submitted to the Programme Coordinator Office must be placed into the assignment box located outside the office.

Office Use Only	
Signature:	
Date:	
Penalty Applied (if applicable):	

Configuration Manual

Anthony Effiok
x19108788

1 Introduction

This configuration manual aims to provide all information in regards to the system configuration, the programming language used and the complete program code used for the research in the project: “NiohSign: A Siamese neural network approach for Signature Authentication”

2 System Configuration

2.1 Hardware

- OS: Mac OS
- RAM: 8GB
- Processor: Inter core i5
- Hard Disk: 250GB

2.2 Software

- Jupyter Notebook - Python

3 Project Development

The data required for the project is a signature dataset known as BHSig260 ¹. This data set comprises of signatures appended by 260 individuals of which 100 are in Bengali and 160 are in Hindi. The signatures appended in Hindi are used to test the performance of the model created in this project.

Figure 1 shows the importation of required libraries for the execution of the project namely the keras libraries.

¹ <https://goo.gl/9QfByd>


```

jupyter Complete NiohSign Last Checkpoint: 3 hours ago (autosaved)
File Edit View Insert Cell Kernel Widgets Help Not Trusted Python 3

In [89]: orig_train, orig_val, orig_test = orig_groups[:120], orig_groups[120:140], orig_groups[140:]
         forg_train, forg_val, forg_test = forg_groups[:120], forg_groups[120:140], forg_groups[140:]

In [90]: # Delete unnecessary variables
         del orig_groups, forg_groups

In [91]: # All the images will be converted to the same size before processing
         img_h, img_w = 299, 299

In [92]: def visualize_sample_signature():
         '''Function to randomly select a signature from train set and
         print two genuine copies and one forged copy'''
         fig, (ax1, ax2, ax3) = plt.subplots(1, 3, figsize = (10, 10))
         k = np.random.randint(len(orig_train))
         orig_img_names = random.sample(orig_train[k], 2)
         forg_img_name = random.sample(forg_train[k], 1)
         orig_img1 = cv2.imread(orig_img_names[0], 0)
         orig_img2 = cv2.imread(orig_img_names[1], 0)
         forg_img = plt.imread(forg_img_name[0], 0)
         orig_img1 = cv2.resize(orig_img1, (img_w, img_h))
         orig_img2 = cv2.resize(orig_img2, (img_w, img_h))
         forg_img = cv2.resize(forg_img, (img_w, img_h))

         ax1.imshow(orig_img1, cmap = 'gray')
         ax2.imshow(orig_img2, cmap = 'gray')
         ax3.imshow(forg_img, cmap = 'gray')

         ax1.set_title('Genuine Copy')
         ax1.axis('off')
         ax2.set_title('Genuine Copy')
         ax2.axis('off')
         ax3.set_title('Forged Copy')
         ax3.axis('off')

```

Fig 3: Further Data exploration

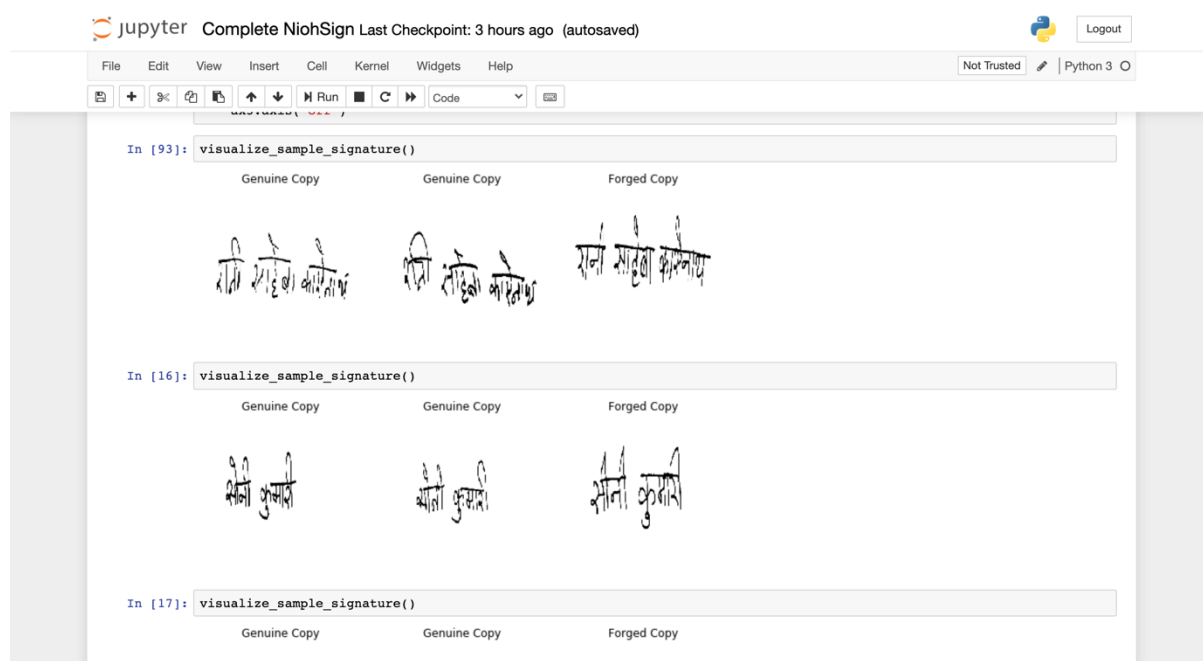


Fig 4: Signature Visualization

Figure 5 shows the generation of the signatures as inputs for the model by splitting them into pairwise of genuine – genuine and genuine- forged signatures.

```

In [94]: def generate_batch(orig_groups, forg_groups, batch_size = 32):
    '''Function to generate a batch of data with batch size number of data points
    Half of the data points will be Genuine-Genuine pairs and half will be Genuine-Forged pairs'''
    while True:
        orig_pairs = []
        forg_pairs = []
        gen_gen_labels = []
        gen_for_labels = []
        all_pairs = []
        all_labels = []

        # Here we create pairs of Genuine-Genuine image names and Genuine-Forged image names
        # For every person we have 24 genuine signatures, hence we have
        # 24 choose 2 = 276 Genuine-Genuine image pairs for one person.
        # To make Genuine-Forged pairs, we pair every Genuine signature of a person
        # with 12 randomly sampled Forged signatures of the same person.
        # Thus we make 24 * 12 = 300 Genuine-Forged image pairs for one person.
        # In all we have 120 person's data in the training data.
        # Total no. of Genuine-Genuine pairs = 120 * 276 = 33120
        # Total number of Genuine-Forged pairs = 120 * 300 = 36000
        # Total no. of data points = 33120 + 36000 = 69120
        for orig, forg in zip(orig_groups, forg_groups):
            orig_pairs.extend(list(itertools.combinations(orig, 2)))
            for i in range(len(forg)):
                forg_pairs.extend(list(itertools.product(orig[i:i+1], random.sample(forg, 12))))

        # Label for Genuine-Genuine pairs is 1
        # Label for Genuine-Forged pairs is 0
        gen_gen_labels = [1]*len(orig_pairs)
        gen_for_labels = [0]*len(forg_pairs)

        # Concatenate all the pairs together along with their labels and shuffle them
        all_pairs = orig_pairs + forg_pairs
        all_labels = gen_gen_labels + gen_for_labels
        del orig_pairs, forg_pairs, gen_gen_labels, gen_for_labels
        all_pairs, all_labels = shuffle(all_pairs, all_labels)

```

Fig 5: Pair wise Split

Figure 6, 7, 8 & 9 represent the InceptionResNetV2 network created for the Siamese network

```

In [99]: # we reduce # filters by factor of 8 compared to original inception-v4
nb_filters_reduction_factor = 8

def inception_resnet_v2_stem(x):
    # in original inception-resnet-v2, conv stride is 2
    x = Convolution2D(32//nb_filters_reduction_factor, 3, 3, subsample=(1, 1), activation='relu',
        init='he_normal', border_mode='valid', dim_ordering='tf')(x)
    x = Convolution2D(32//nb_filters_reduction_factor, 3, 3, subsample=(1, 1), activation='relu',
        init='he_normal', border_mode='valid', dim_ordering='tf')(x)
    x = Convolution2D(64//nb_filters_reduction_factor, 3, 3, subsample=(1, 1), activation='relu',
        init='he_normal', border_mode='same', dim_ordering='tf')(x)

    # in original inception-resnet-v2, stride is 2
    a = MaxPooling2D((3, 3), strides=(1, 1), border_mode='valid', dim_ordering='tf')(x)
    # in original inception-resnet-v2, conv stride is 2
    b = Convolution2D(96//nb_filters_reduction_factor, 3, 3, subsample=(1, 1), activation='relu',
        init='he_normal', border_mode='valid', dim_ordering='tf')(x)
    x = concatenate([a, b], axis=-1)

    a = Convolution2D(64//nb_filters_reduction_factor, 1, 1, subsample=(1, 1), activation='relu',
        init='he_normal', border_mode='same', dim_ordering='tf')(x)
    a = Convolution2D(96//nb_filters_reduction_factor, 3, 3, subsample=(1, 1), activation='relu',
        init='he_normal', border_mode='valid', dim_ordering='tf')(a)
    b = Convolution2D(64//nb_filters_reduction_factor, 1, 1, subsample=(1, 1), activation='relu',
        init='he_normal', border_mode='same', dim_ordering='tf')(x)
    b = Convolution2D(64//nb_filters_reduction_factor, 7, 1, subsample=(1, 1), activation='relu',
        init='he_normal', border_mode='same', dim_ordering='tf')(b)
    b = Convolution2D(64//nb_filters_reduction_factor, 1, 7, subsample=(1, 1), activation='relu',
        init='he_normal', border_mode='same', dim_ordering='tf')(b)
    b = Convolution2D(96//nb_filters_reduction_factor, 3, 3, subsample=(1, 1), activation='relu',
        init='he_normal', border_mode='valid', dim_ordering='tf')(b)
    x = concatenate([a, b], axis=-1)

    # in original inception-resnet-v2, conv stride should be 2

```

Fig 6: InceptionResNetV2 Network (1)



```
def inception_resnet_v2_reduction_B(x):
    a = MaxPooling2D((3, 3), strides=(2, 2), border_mode='valid', dim_ordering='tf')(x)
    b = Convolution2D(256//nb_filters_reduction_factor, 1, 1, subsample=(1, 1), activation='relu',
        init='he_normal', border_mode='same', dim_ordering='tf')(a)
    b = Convolution2D(288//nb_filters_reduction_factor, 3, 3, subsample=(2, 2), activation='relu',
        init='he_normal', border_mode='valid', dim_ordering='tf')(b)
    c = Convolution2D(256//nb_filters_reduction_factor, 1, 1, subsample=(1, 1), activation='relu',
        init='he_normal', border_mode='same', dim_ordering='tf')(x)
    c = Convolution2D(288//nb_filters_reduction_factor, 3, 3, subsample=(2, 2), activation='relu',
        init='he_normal', border_mode='valid', dim_ordering='tf')(c)
    d = Convolution2D(256//nb_filters_reduction_factor, 1, 1, subsample=(1, 1), activation='relu',
        init='he_normal', border_mode='same', dim_ordering='tf')(x)
    d = Convolution2D(288//nb_filters_reduction_factor, 3, 3, subsample=(1, 1), activation='relu',
        init='he_normal', border_mode='same', dim_ordering='tf')(d)
    d = Convolution2D(320//nb_filters_reduction_factor, 3, 3, subsample=(2, 2), activation='relu',
        init='he_normal', border_mode='valid', dim_ordering='tf')(d)

    x = concatenate([a, b, c, d],axis=-1)

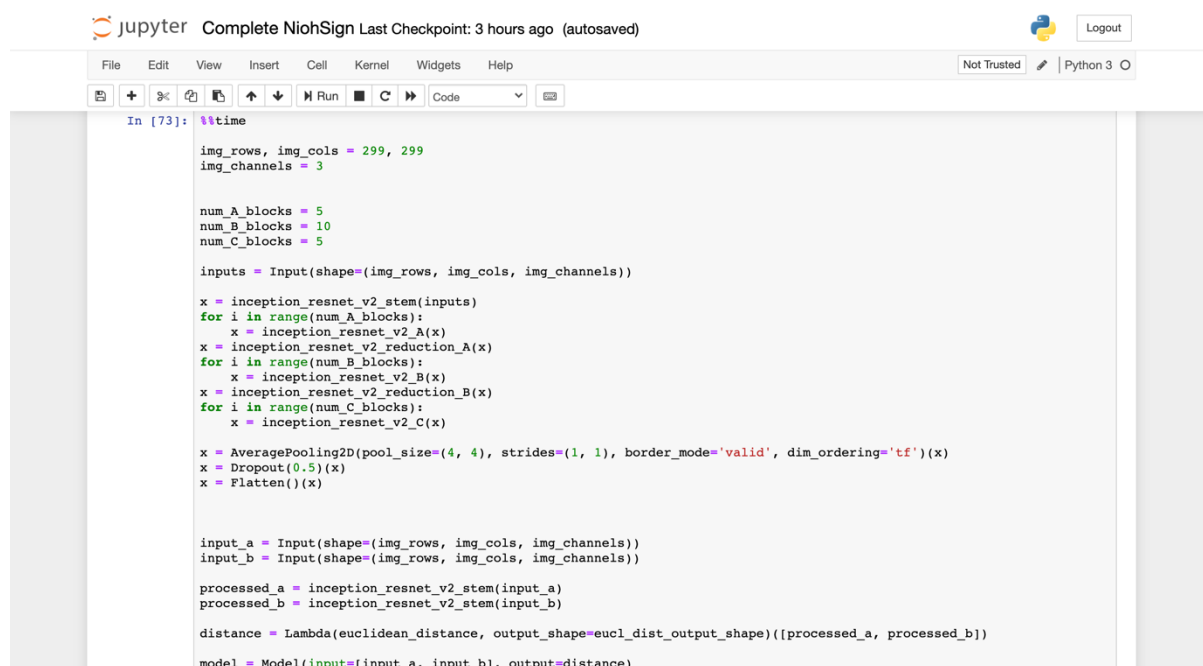
    return x

def inception_resnet_v2_C(x):
    shortcut = x

    a = Convolution2D(192//nb_filters_reduction_factor, 1, 1, subsample=(1, 1), activation='relu',
        init='he_normal', border_mode='same', dim_ordering='tf')(x)
    b = Convolution2D(192//nb_filters_reduction_factor, 1, 1, subsample=(1, 1), activation='relu',
        init='he_normal', border_mode='same', dim_ordering='tf')(x)
    b = Convolution2D(224//nb_filters_reduction_factor, 1, 3, subsample=(1, 1), activation='relu',
        init='he_normal', border_mode='same', dim_ordering='tf')(b)
    b = Convolution2D(256//nb_filters_reduction_factor, 3, 1, subsample=(1, 1), activation='relu',
        init='he_normal', border_mode='same', dim_ordering='tf')(b)
```

Fig 9: InceptionResNetV2 Network (4)

Fig 10 represents the steps taken to create the Siamese Network from the above base network.



```
In [73]: %%time

img_rows, img_cols = 299, 299
img_channels = 3

num_A_blocks = 5
num_B_blocks = 10
num_C_blocks = 5

inputs = Input(shape=(img_rows, img_cols, img_channels))

x = inception_resnet_v2_stem(inputs)
for i in range(num_A_blocks):
    x = inception_resnet_v2_A(x)
x = inception_resnet_v2_reduction_A(x)
for i in range(num_B_blocks):
    x = inception_resnet_v2_B(x)
x = inception_resnet_v2_reduction_B(x)
for i in range(num_C_blocks):
    x = inception_resnet_v2_C(x)

x = AveragePooling2D(pool_size=(4, 4), strides=(1, 1), border_mode='valid', dim_ordering='tf')(x)
x = Dropout(0.5)(x)
x = Flatten()(x)

input_a = Input(shape=(img_rows, img_cols, img_channels))
input_b = Input(shape=(img_rows, img_cols, img_channels))

processed_a = inception_resnet_v2_stem(input_a)
processed_b = inception_resnet_v2_stem(input_b)

distance = Lambda(euclidean_distance, output_shape=eucl_dist_output_shape)([processed_a, processed_b])

model = Model(input=[input_a, input_b], output=distance)
```

Fig 10: Creation of the Siamese network

Figure 11 represents the summary of the created model

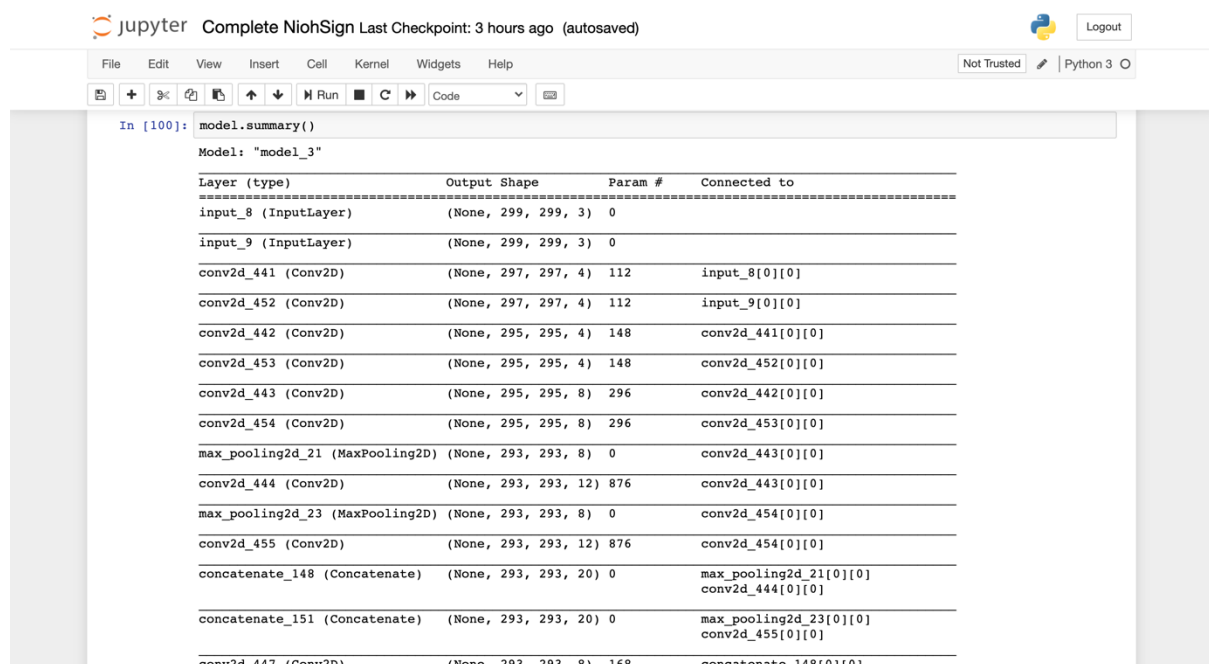


Fig 11: Model Summary

Figure 12 shows the steps taken to fit and run the model

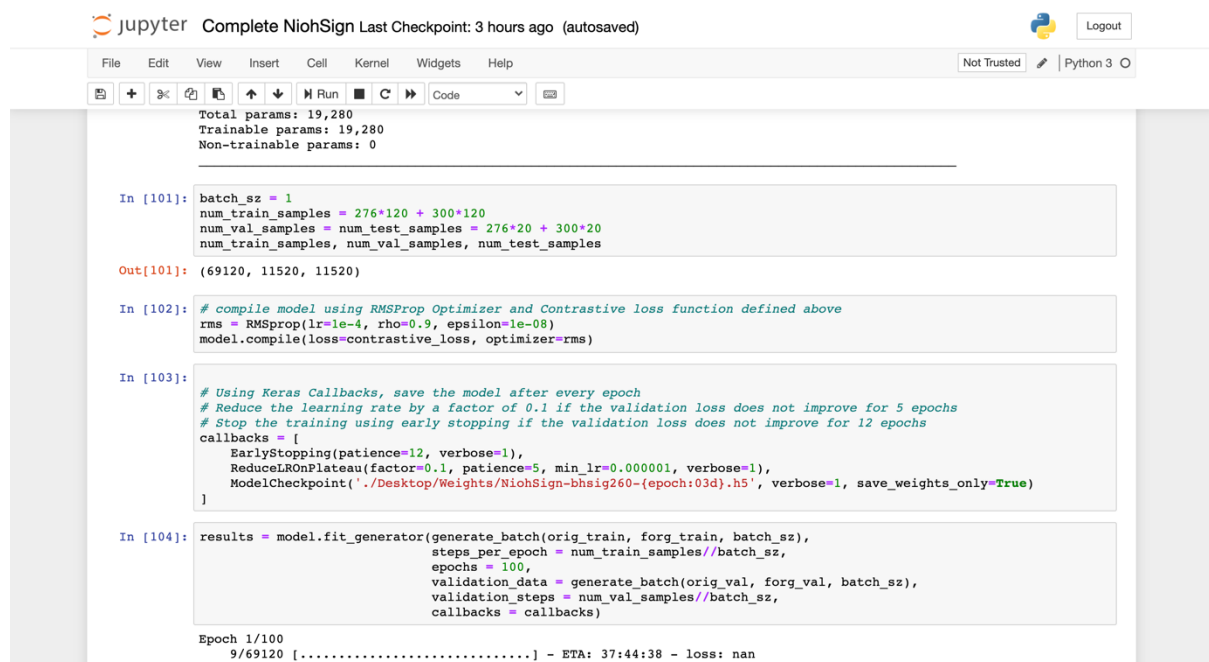


Fig 12: Running the Model

Figure 13 shows the steps to load the weights after running the model and the creation of the function for the accuracy and the threshold that will be used for the signatures.



The Jupyter Notebook interface shows a Python script for calculating ROC accuracy and loading model weights. The script defines a function `compute_accuracy_roc` that iterates over a range of thresholds to find the one that maximizes accuracy. It then loads the model weights from a file named `Weights/NiohSign-bhsig260-(epoch:03d).h5`. Finally, it generates a batch of test data and begins a loop to process each sample.

```

In [105]: def compute_accuracy_roc(predictions, labels):
            '''Compute ROC accuracy with a range of thresholds on distances.
            ...
            dmax = np.max(predictions)
            dmin = np.min(predictions)
            nsame = np.sum(labels == 1)
            ndiff = np.sum(labels == 0)

            step = 0.01
            max_acc = 0
            best_thresh = -1

            for d in np.arange(dmin, dmax+step, step):
                idx1 = predictions.ravel() <= d
                idx2 = predictions.ravel() > d

                tpr = float(np.sum(labels[idx1] == 1)) / nsame
                tnr = float(np.sum(labels[idx2] == 0)) / ndiff
                acc = 0.5 * (tpr + tnr)
                # print ('ROC', acc, tpr, tnr)

                if (acc > max_acc):
                    max_acc, best_thresh = acc, d

            return max_acc, best_thresh

Load the weights from the epoch which gave the best validation accuracy

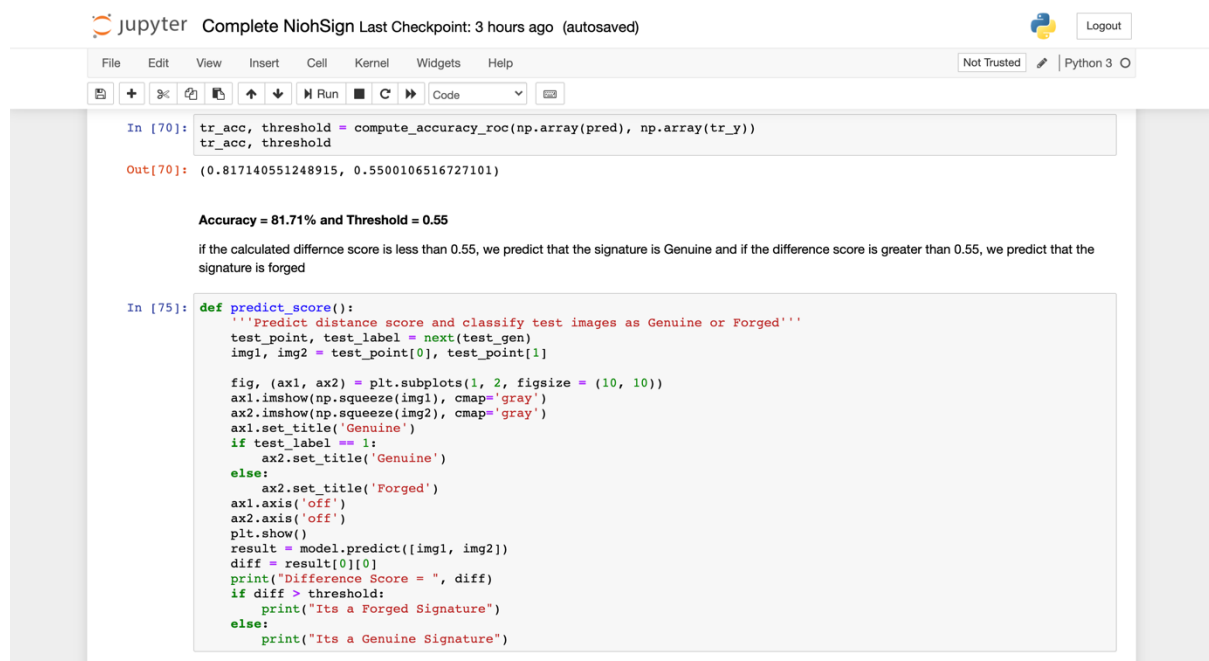
In [ ]: model.load_weights('./Weights/NiohSign-bhsig260-(epoch:03d).h5')

In [69]: test_gen = generate_batch(orig_test, forg_test, 1)
         pred, tr_y = [], []
         for i in range(num_test_samples):
             (img1, img2), label = next(test_gen)
             ...

```

Fig 13: Threshold and accuracy calculation

Figure 14 shows accuracy and threshold along with the function for the prediction of the signatures.



The Jupyter Notebook interface shows the execution of the prediction function. It calls `compute_accuracy_roc` with the model's predictions and true labels, resulting in an accuracy of 0.8171 and a threshold of 0.5501. Below this, a function `predict_score` is defined to take test images and labels, display them side-by-side, and print the predicted score and the difference score. The function uses `plt.imshow` to show the images and `plt.title` to label them as 'Genuine' or 'Forged'.

```

In [70]: tr_acc, threshold = compute_accuracy_roc(np.array(pred), np.array(tr_y))
         tr_acc, threshold

Out[70]: (0.817140551248915, 0.5500106516727101)

Accuracy = 81.71% and Threshold = 0.55

if the calculated difference score is less than 0.55, we predict that the signature is Genuine and if the difference score is greater than 0.55, we predict that the signature is forged

In [75]: def predict_score():
            '''Predict distance score and classify test images as Genuine or Forged'''
            test_point, test_label = next(test_gen)
            img1, img2 = test_point[0], test_point[1]

            fig, (ax1, ax2) = plt.subplots(1, 2, figsize = (10, 10))
            ax1.imshow(np.squeeze(img1), cmap='gray')
            ax2.imshow(np.squeeze(img2), cmap='gray')
            ax1.set_title('Genuine')
            if test_label == 1:
                ax2.set_title('Genuine')
            else:
                ax2.set_title('Forged')
            ax1.axis('off')
            ax2.axis('off')
            plt.show()
            result = model.predict([img1, img2])
            diff = result[0][0]
            print("Difference Score = ", diff)
            if diff > threshold:
                print("Its a Forged Signature")
            else:
                print("Its a Genuine Signature")

```

Fig 14: Prediction function

Figure 15 shows the output of a prediction which is representative of the ability of the network to classify signatures as authentic or forged



Fig 15: Final Outputs