

Configuration Manual

MSc Data Analytics Research Project

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MSc Project Submission Sheet



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Configuration Manual

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1 Introduction

This Configuration Manual lists all prerequisites needed to duplicate the studies and their effects on a specific setting. A glimpse of the source for Data Importing & Classifying data as per categories and after that images are augmented, and prediction algorithms are built for detection of class.

The report is organized as follows, with details relating to environment configuration provided in Section 2. Information about data gathering is detailed in Section 3. Data Classification is done in Section 4. Image Segmentation is included in Section 5. In section 6, the Image Augmentation is described. Details about models that were created and tested are provided in Section 7. How the results are calculated and shown is described in Section 8.

2 System Requirements

The specific needs for hardware and software to put the research into use are detailed in this section.

2.1 Hardware Requirements

2.1 Hardware Requirements

The code execution took place on my HP Spectre x360 laptop, featuring the hardware specifications detailed in Figure 1:

- Processor: Intel Core i7 12th Gen
- Operating System: 64-bit
- RAM: 16GB
- Storage: 256GB SSD
- Display: 14-inch

It is anticipated that the code is compatible with a configuration comprising an i3 processor, 8GB RAM, and a 256GB SSD. However, it is pertinent to mention that no specific testing has been conducted on this setup.

System > About			
Mrun HP Sp	ali ectre x360 2-in-1 Lap	ptop 14-ef0xxx	Rename this PC
(j	Device specificat	tions	Сору ^
	Device name	Mrunali	
	Processor	12th Gen Intel(R) Core(TM) i7-1255U 1.70 GHz	
	Installed RAM 16.0 GB (15.7 GB usable)		
	Device ID FC951C9C-C04E-4574-A623-D2A9DFDF862A		
	Product ID 00342-42617-76026-AAOEM		
	System type	64-bit operating system, x64-based processor	
	Pen and touch	Pen and touch support with 256 touch points	

Figure 1. Hardware Requirement

2.2 Software Requirements

- \Box Anaconda 3 (Version 4.8.0)
- \Box Jupyter Notebook (Version 6.0.3)
- \Box Python (Version 3.7.6)

2.3 Code Execution

The code can be run in jupyter notebook. The Jupyter Notebook comes with Anaconda 3, running the Jupyter Notebook from startup. This will open Jupyter Notebook in the web browser. The web browser will show the folder structure of the system and move to the folder where the code file is located. Open the code file from the folder and to run the code, go to the Kernel menu and run all cells.

3 Data Collection

The dataset is taken from the Kaggle public repository from the link https://www.kaggle.com/competitions/diabetic-retinopathy-detection/data. a large set of high-resolution retina images taken under a variety of imaging conditions. A left and right field is provided for every subject. Images are labeled with a subject ID as well as either left or right (e.g. 1_left.jpeg is the left eye of the patient id 1)

4 Data Classification

Figure 2 lists every Python library necessary to complete the project.

import numpy as np import pandas as pd import matplotlib.pyplot as plt import seaborn as sn import warnings import os import shutil warnings.filterwarnings('ignore') import glob import cv2 from PIL import ImageFile import tensorflow import keras from tensorflow.keras.preprocessing.image import ImageDataGenerator from tensorflow.keras.models import Sequential, Model from tensorflow.keras.layers import GlobalAveragePooling2D, Conv2D, Dense, Flatten, Layer, Input, Dropout, BatchNormalization from tensorflow.keras.layers import Attention, MultiHeadAttention, LayerNormalization, Add, Multiply, MaxPooling2D from keras.applications.inception v3 import InceptionV3 from keras.applications.resnet v2 import ResNet50V2 from keras.applications.efficientnet v2 import EfficientNetV2L

Figure 2: Necessary Python libraries

Figure 3 represents the block of code to import the training videos and check for the data.

```
#Importing Data Set CSV file
trainLabels=pd.read_csv(r'trainLabels.csv')
#Dataset Information
trainLabels.head(10)
     image level
 0
    10_left
               0
 1 10_right
               0
 2
    13_left
               0
 3 13_right
               0
    15 left
               1
 4
 5
  15_right
               2
 6
   16_left
               4
  16_right
 7
               4
 8
   17_left
               0
 9 17_right
               1
```

Figure 3: Importing training videos and Checking Data Information

As seen in Figure 4, appending '.jpeg' in the image column to complete the filename.

trainL trainL	abels[<mark>'image'</mark> abels]= tr
	image	level
0	10_left.jpeg	0
1	10_right.jpeg	0
2	13_left.jpeg	0
3	13_right.jpeg	0
4	15_left.jpeg	1
35121	44347_right.jpeg	0
35122	44348_left.jpeg	0
35123	44348_right.jpeg	0
35124	44349_left.jpeg	0
35125	44349_right.jpeg	1

³⁵¹²⁶ rows × 2 columns

Figure 5, the code generates a subset of data based on classes of images and creates a Data folder to classify data based on each category in the folder.



Figure 5: Data Categorization

Figure 6, illustrates the code to classify images into folders based on their category.

```
def make direct(Disease,image):
        global Diseases_path
        if image in Image_Data:
               if os.path.exists(os.path.join(Destination path+Disease)):
                       pass
                else:
                       os.makedirs(Destination_path+Disease)
                       print(image)
               return shutil.copyfile(os.path.join(IMAGE_PATH,image),os.path.join((Destination_path+Disease),image))
                                                                                            #Created Directory for Moderate Diabetic Retinopathy
moderate = [Moderate['image'].apply(lambda x: make_direct(r'\Moderate',x))]
moderate
                                                                                           [5
23
V
                                                                                                         Data\Moderate\15_right.jpeg
None
 #Created Directory for No Diabetic Retinopathy
noDR = [NoDR['image'].apply(lambda x: make_direct(r'\NoDR',x))]
                                                                                                                                         None
None
None
                                                                                             30
42
46
 noDR
                Data\NoDR\10_left.jpeg
 [0]
                                                                                             35005
35018
35019
                                                                                                                                         None
None
None
None
              Data\NoDR\10_right.jpeg
                Data\NoDR\13 left.jpeg
  2
                                                                                              35034
              Data\NoDR\13_right.jpeg
                                                                                              35063
                                                                                                                                         Non
                                                                                             Name: image, Length: 5292, dtype: object]
  8
                Data\NoDR\17_left.jpeg
                           ...
                                                                                                        Directory for Severe Diabetic Retinopathy
[Severe['image'].apply(lambda x: make_direct(r'\Severe',x))]
  35120
                                         None
  35121
                                         None
                                                                                           [90
91
134
135
283
  35122
                                         None
  35123
                                         None
                                         None
                                                                                                           Data\Severe\163_left
Data\Severe\163_right
  35124
                                                                                                                                 ight.
  Name: image, Length: 25810, dtype: object]
                                                                                             34915
                                                                                                                                       None
 #Created Directory for Mild Diabetic Retinopathy
mild = [Mild['image'].apply(lambda x: make_direct(r'\Mild',x))]

        34915
        Nome

        34961
        None

        34994
        None

        34995
        None

        34997
        None

        Name: image, Length: 873, dtype: object]

 mild
                Data\Mild\15_left.jpeg
 [4
  .
9
              Data\Mild\17_right.jpeg
                                                                                            #Created Directory for Proliferative Diabetic Retinopathy
proliferative = [Proliferative['image'].apply(lambda x: make_direct(r'\Proliferative',x))]
  22
                                         None
                                                                                            proliferative
                                         None
   28
                                                                                                       ative
Data\Proliferative\16_left.jpeg
Data\Proliferative\16_right.jpeg
Data\Proliferative\217_left.jpeg
Data\Proliferative\217_right.jpeg
None
              Data\Mild\114_left.jpeg
  102
                                                                                            [6
                           ...
                                                                                             184
  35098
                                         None
                                                                                             185
252
  35099
                                         None
  35105
                                         None
                                                                                             34914
35046
                                                                                                                                                 None
  35112
                                         None
                                                                                              35040
   35125
                                         None
                                                                                              35048
  Name: image, Length: 2443, dtype: object]
                                                                                              35049
                                                                                             Name: image, Length: 708, dtype: object]
```

Figure 6: Data Categorization

Figure 7 illustrates the code to update the data path and set image size and categories.

```
# PATH Initilization
DATA_PATH=Destination_path
# Initilizing parameters related to images
IMAGE_WIDTH= 256
IMAGE_HEIGHT= 256
path_noDR = './Data/NoDR/'
path_mild = './Data/Moderate/'
path_moderate = './Data/Moderate/'
path_severe = './Data/Severe/'
path_proliferative = './Data/Proliferative/'
```

```
categories = ['NoDR', 'Mild', 'Moderate', 'Severe', 'Proliferative']
print(categories)
```

['NoDR', 'Mild', 'Moderate', 'Severe', 'Proliferative']

Figure 7: Data Path

Figure 8 and Figure 9, illustrate the check count of images in each category and generate a bar plot of the value counts.



categories = ['No DR', 'Mild', 'Moderate', 'Severe', 'Proliferative']
values = [count_nodr, count_mild, count_moderate, count_sevre, count_proliferative]

```
# Plotting the pie chart
plt.figure(figsize=(8, 8))
plt.pie(values, labels=categories, autopct='%1.1f%%', startangle=140, colors=['#66b3ff',
plt.title('Distribution of Classes in the Dataset (Pie Chart)')
plt.show()
```





Figure 9: Data Classes Pie chart

Figure 10 illustrates the generated list of images for each class.

```
nodr = glob.glob(path_noDR+ "*.jpeg")
# Print out the first 5 file names to verify we're in the right folder.
print ("Total of %d images.\nFirst 5 filenames:" % len(nodr))
print ('\n'.join(nodr[:5]))
Total of 12387 images.
First 5 filenames:
./Data/NoDR\10003_left.jpeg
./Data/NoDR\10003_right.jpeg
./Data/NoDR\10007_left.jpeg
./Data/NoDR\10007_right.jpeg
./Data/NoDR\10009_left.jpeg
mild = glob.glob(path_mild + "*.jpeg")
# Print out the first 5 file names to verify we're in the right folder.
print ("Total of %d images.\nFirst 5 filenames:" % len(mild))
print ('\n'.join(mild[:5]))
Total of 1130 images.
First 5 filenames:
./Data/Mild\10030 left.jpeg
./Data/Mild\10030_right.jpeg
./Data/Mild\10085_left.jpeg
./Data/Mild\10085_right.jpeg
./Data/Mild\10150_right.jpeg
moderate = glob.glob(path moderate + "*.jpeg")
# Print out the first 5 file names to verify we're in the right folder.
print ("Total of %d images.\nFirst 5 filenames:" % len(moderate))
print ('\n'.join(moderate[:5]))
Total of 2537 images.
First 5 filenames:
./Data/Moderate\1002 right.jpeg
./Data/Moderate\10043_left.jpeg
./Data/Moderate\10043_right.jpeg
./Data/Moderate\10109_left.jpeg
./Data/Moderate\10109_right.jpeg
severe = glob.glob(path severe + "*.jpeg")
# Print out the first 5 file names to verify we're in the right folder.
print ("Total of %d images.\nFirst 5 filenames:" % len(severe))
print ('\n'.join(severe[:5]))
Total of 424 images.
First 5 filenames:
./Data/Severe\1002_left.jpeg
./Data/Severe\10047_left.jpeg
./Data/Severe\1008 left.jpeg
./Data/Severe\1008_right.jpeg
./Data/Severe\10125_left.jpeg
proliferative = glob.glob(path_proliferative + "*.jpeg")
# Print out the first 5 file names to verify we're in the right folder.
print ("Total of %d images.\nFirst 5 filenames:" % len(proliferative))
print ('\n'.join(proliferative[:5]))
Total of 324 images.
First 5 filenames:
./Data/Proliferative\10017_left.jpeg
./Data/Proliferative\10017_right.jpeg
./Data/Proliferative\10047_right.jpeg
./Data/Proliferative\10193_right.jpeg
./Data/Proliferative\10312_left.jpeg
```

Figure 10: Images in each class

5 Image Segmentation

Figure 11 illustrates the read the image and shows the image shape and plot image.



Figure 11: Read Image

Figures 12 show the code used to read the image in gray scale.

```
img1_gray = cv2.cvtColor(img1, cv2.COLOR_RGB2GRAY)
plot_image(img1_gray)
width, height = img1_gray.shape
print(f'Width Grayscale = {width}')
print(f'Height Grayscale = {height}')
print(f'Image Shape Grayscale {img1_gray.shape}')
```

```
Width Grayscale = 3168
Height Grayscale = 4752
Image Shape Grayscale (3168, 4752)
```



Figure 12: Read GrayScale Image



Figure 13 illustrates the code to generate adaptive thresholds of the images and display the contours.

Figure 13: Adaptive threshold and contouring

1000

6 Image Augmentation

Figure 14 illustrates the code to use ImageDataGenertor to generate augmented images for the deep learning models.



20 40 60 80 0 20 40 60 80 0 20 40 60 80 0 20 40 60

Figure 14: Image Data Generator

Figures 15 show the code to create data with different height and width shifts to check for appropriate size.

```
gen = ImageDataGenerator(rescale=1./255, width_shift_range=0.2, height_shift_range=0.3)
train = gen.flow_from_directory(directory=DATA_PATH, target_size=(img_size,img_size), class_mode='sparse')
augmented_images = [train[0][0][0] for i in range(4)]
plotImages(augmented_images)
```

Found 16802 images belonging to 5 classes.



Figure 15: Image Data Generator

Figure 16 shows the code to create data by rescaling the images.

gen = ImageDataGenerator(rescale=1./255)
train = gen.flow_from_directory(directory=DATA_PATH, target_size=(img_size,img_size))
augmented_images = [train[0][0][0] for i in range(4)]
plotImages(augmented_images)

Found 16802 images belonging to 5 classes.



Figure 16: Image Data Generator

Figure 17 illustrates the code to data and validation split to generate train and test data.

Found 1678 images belonging to 5 classes.

```
input_shape = (img_size, img_size, 3)
input_shape
```

(100, 100, 3)

Figure 17: Image Data Generator

7 Machine Learning Models

7.1 Transformer CNN

```
flatten = Flatten()(decoder)
x = Dense(5, activation='relu')(flatten)
model = Model(inputs=input, outputs=x)
model.summary()
model.compile(optimizer = 'adam', loss= 'categorical_crossentropy', metrics = ['accuracy'])
```

Model: "model 1"

Layer (type)	Output Shape	Param #	Connected to
input_3 (InputLayer)	[(None, 100, 100, 3)]	0	[]
layer_normalization_6 (Lay erNormalization)	(None, 100, 100, 3)	6	['input_3[0][0]']
dense_11 (Dense)	(None, 100, 100, 128)	512	['layer_normalization_6[0][0]]
dense_12 (Dense)	(None, 100, 100, 3)	387	['dense_11[0][0]']
add_7 (Add)	(None, 100, 100, 3)	0	['layer_normalization_6[0][0] , 'dense_12[0][0]']
layer_normalization_7 (Lay erNormalization)	(None, 100, 100, 3)	6	['add_7[0][0]']

model.fit(train)

473/473 [======] - 2389s 5s/step - loss: 2.7787 - accuracy: 0.7073

<keras.src.callbacks.History at 0x1e18df84910>

Figure 18: Implementation of Transformer CNN

7.2 ResNet 50

model = ResNet50V2(include_top=False, weights='imagenet', input_shape=input_shape)
model.summary()

Model: "resnet50v2"

Layer (type)	Output Shape	Param #	Connected to
input_5 (InputLayer)	[(None, 100, 100, 3)]	0	[]
conv1_pad (ZeroPadding2D)	(None, 106, 106, 3)	0	['input_5[0][0]']
<pre>conv1_conv (Conv2D)</pre>	(None, 50, 50, 64)	9472	['conv1_pad[0][0]']
pool1_pad (ZeroPadding2D)	(None, 52, 52, 64)	0	['conv1_conv[0][0]']
<pre>pool1_pool (MaxPooling2D)</pre>	(None, 25, 25, 64)	0	['pool1_pad[0][0]']
conv2_block1_preact_bn (Ba tchNormalization)	(None, 25, 25, 64)	256	['pool1_pool[0][0]']
conv2_block1_preact_relu (Activation)	(None, 25, 25, 64)	0	['conv2_block1_preact_bn[0][0] ']

```
x = model.output
x = Flatten()(x)
x = Dense(512, activation='tanh')(x)
x = Dropout(0.5)(x)
x = Dense(128, activation='tanh')(x)
x = Dropout(0.01)(x)
x = Dense(64, activation='tanh')(x)
x = Dropout(0.05)(x)
output_layer = Dense(1, activation='sigmoid')(x)
```

model = Model(inputs=model.input, outputs=output_layer)

model.compile(optimizer = 'adam', loss= 'binary_crossentropy', metrics = ['accuracy'])
model.summary()

Model: "model_2"

Layer (type)	Output Shape	Param #	Connected to
input_5 (InputLayer)	[(None, 100, 100, 3)]	0	[]
conv1_pad (ZeroPadding2D)	(None, 106, 106, 3)	0	['input_5[0][0]']
conv1_conv (Conv2D)	(None, 50, 50, 64)	9472	['conv1_pad[0][0]']
pool1_pad (ZeroPadding2D)	(None, 52, 52, 64)	0	['conv1_conv[0][0]']
<pre>pool1_pool (MaxPooling2D)</pre>	(None, 25, 25, 64)	0	['pool1_pad[0][0]']
conv2_block1_preact_bn (Ba tchNormalization)	(None, 25, 25, 64)	256	['pool1_pool[0][0]']
conv2_block1_preact_relu (Activation)	(None, 25, 25, 64)	0	['conv2_block1_preact_bn[0][0] ']

model.fit(train, batch_size=500)

<keras.src.callbacks.History at 0x1e1815f0be0>

Figure 19: Implementation of ResNet50

7.3 InceptionNet

```
base_model = InceptionV3(weights='imagenet', include_top=False)
base_model.summary
```

<bound method Model.summary of <keras.src.engine.functional.Functional object at 0x000001E185E74D60>>

```
x = base_model.output
x = GlobalAveragePooling2D()(x)
x = Dense(512, activation='relu')(x)
x = Dropout(0.15)(x)
x = Dense(128, activation='sigmoid')(x)
x = Dropout(0.05)(x)
x = Dense(8, activation='sigmoid')(x)
predictions = Dense(5, activation='sigmoid')(x)
```

model = Model(inputs=base_model.input, outputs=predictions)

```
model.compile(optimizer = 'adam', loss= 'categorical_crossentropy', metrics = ['accuracy'])
model.summary()
```

Model: "model_3"

Layer (type)	Output Shape	Param #	Connected to
input_6 (InputLayer)	[(None, None, None, 3)]	0	[]
conv2d (Conv2D)	(None, None, None, 32)	864	['input_6[0][0]']
batch_normalization (Batch Normalization)	(None, None, None, 32)	96	['conv2d[0][0]']
activation (Activation)	(None, None, None, 32)	0	['batch_normalization[0][0]']
conv2d_1 (Conv2D)	(None, None, None, 32)	9216	['activation[0][0]']
<pre>batch_normalization_1 (Bat chNormalization)</pre>	(None, None, None, 32)	96	['conv2d_1[0][0]']
activation_1 (Activation)	(None, None, None, 32)	0	['batch_normalization_1[0][0]'

model.fit(train, batch_size=5000)

473/473 [=======] - 3399s 7s/step - loss: 0.9909 - accuracy: 0.7331

<keras.src.callbacks.History at 0x1e180f66f40>

Figure 20: Implementation of InceptionNet V3

7.4 EfficientNet

```
import ssl
ssl._create_default_https_context = ssl._create_unverified_context
```

```
base_model = EfficientNetV2L(weights='imagenet', include_top=False)
base_model.summary
```

<bound method Model.summary of <keras.src.engine.functional.Functional object at 0x000001E200EFA850>>

```
x = base_model.output
x = GlobalAveragePooling2D()(x)
x = Dropout(0.05)(x)
x = Dense(8, activation='sigmoid')(x)
predictions = Dense(5, activation='softmax')(x)
```

```
model = Model(inputs=base_model.input, outputs=predictions)
```

```
model.compile(optimizer = 'adam', loss= 'categorical_crossentropy', metrics = ['accuracy'])
model.summary()
```

Model: "model_4"

Layer (type)	Output Shape	Param #	Connected to
input_7 (InputLayer)	[(None, None, None, 3)]	0	[]
rescaling (Rescaling)	(None, None, None, 3)	0	['input_7[0][0]']
stem_conv (Conv2D)	(None, None, None, 32)	864	['rescaling[0][0]']
stem_bn (BatchNormalizatio n)	(None, None, None, 32)	128	['stem_conv[0][0]']
stem_activation (Activatio n)	(None, None, None, 32)	0	['stem_bn[0][0]']
block1a_project_conv (Conv 2D)	(None, None, None, 32)	9216	['stem_activation[0][0]']
blackt and be (Databas	(New New 20)	100	[1]]]]]]]]]]]]]]]]]]]]]]]]]]]]]]]]]]]]]

model.fit(train, batch_size=500)

473/473 [=========] - 7400s 15s/step - loss: 0.8902 - accuracy: 0.7359

<keras.src.callbacks.History at 0x1e200ef6160>

Figure 21: Implementation of EfficientNet

References

https://www.kaggle.com/competitions/diabetic-retinopathy-detection/data

OpenCV: OpenCV modules

Keras Applications

ttps://www.picsellia.com/post/are-transformers-replacing-cnns-in-object-detection