

# **Configuration Manual**

MSc Research Project Data Analytics

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#### National College of Ireland

#### **MSc Project Submission Sheet**



Year: 2019-2020

#### **School of Computing**

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Programme: Data Analytics

Module: MSc Research Project

Lecturer: MR. Hicham Rifai Submission

**Due Date:** 17/08/2020

Project Title: Configuration Manual

Word Count: 720

Page Count: 12

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# **Configuration Manual**

### Shovan Roy X18201156

### **1** Introduction

This Configuration manual represents all the necessary requirements and details for the Project "COVID-19 Detection using Deep Learning". The 2<sup>nd</sup> section shows the Hardware requirements. The third section focuses on all the software required to be able to run this project. The 4<sup>th</sup> section Shows where to get the data from. The 5<sup>th</sup> section depicts the library module requirements of the project. The rest of the sections are divided into parts of workflow of the code, their execution, result and their evaluation.

# 2 Hardware Requirements

ASUS ROG 64 bits windows 10 operating system has been used for the project.



**Figure 1 Hardware Requirements** 

### **3** Software Requirements

The Entire Script has been implemented using python 3.8[1]. The Anaconda Navigator[2][3] is required because the whole program has been done in Jupyter Notebook which in included in Anaconda Navigator.

ome	Applications on base (root)	✓ Channels					
vironments	Ô	¢	¢ jupyter	Ô	Ê	¢ IP[y]:	
rning	CMD.exe Prompt	JupyterLab	Notebook	Powershell Prompt	PyCharm	Qt Console	
nmunity	0.1.1 Run a cmd.exe terminal with your current environment from Navigator activated	An extensible environment for interactive and reproducible computing, based on the Jupyter Notebook and Architecture.	6.0.3 Web-based, interactive computing notebook environment. Edit and run human-readable docs while describing the data analysis.	0.0.1 Run a Powershell terminal with your current environment From Navigator activated	2020.2 Full-Featured Python IDE by JetBrains. Supports code completion, linting, debugging, and domain-specific enhancements for web development and data science.	4.7.5 PyQt GUI the supports inline figures, proper multiline editing with syntax highlighting, graphical calitips, and more.	
	Launch	Launch	Launch	Launch	Leunch	Launch	
	Soyder 4.14 Scientific Proto-Development Bruiklonnest, Poverful Syston OE victo advanced delin, interactive testing, debugging and introspection features	Clueviz Glueviz 0.13 Multidimensional dea visualitation across fice. Explore relationships within and among related datasets.	Crange 3 3.48 Concorrect back data mixing framework. Data visualization and data analysis for noisic and example. Thereative workflows with a large toolbox.	Riculo 1.48 Ast of integrated tool deligned to help public more productive with R. Includes R essentials and notebook.			
	Launch	install	install	Install			
umentation							

Figure 2: Anaconda Navigator



**Figure 3: Jupyter Notebook** 

### **4** Data Requirements

The Data required for the project can be Downloaded from the Github[4] and contains no personal data and is in accordance with the GDPR Guidelines.

→ C (	github.com/ieee8023/covid-chestxray-da	taset		* 🖬 🖬 🦉 🐂 🖾 🗇 🛪	- ⊒ <u>(</u>		
Search or	r jump to Pull rec	uests Issues Marketplace Explore		Ċ	+• 🗳		
ieee8023	3 / covid-chestxray-dataset			O Watch ▼ 145 ☆ Star 2.3k ♥ For     For     Star 2.3k ♥ F	k 766		
<> Code	() Issues 27 Il Pull requests 1		ecurity 🗠 Insights				
	🐉 master 👻 🐉 25 branches		Go to file Add file ▼ 👱 Code →	About			
	bganglia and ieee8023 More	RT-PCR annotations	✓ 6433253 2 days ago ⑦ 680 commits	We are building an open database of COVID-19 cases with chest X-ray or CT images			
	.github/workflows	Update tests.yml	3 months ago	covid-19 deen-learning			
	annotations	Update README.md	3 months ago	computer-vision dataset xray			
	docs	add share image	5 months ago	computed-tomography			
	images	Add Radiopaedia images	7 days ago	🖾 Readme			
	scripts	docstring for cache browsing script	6 days ago				
	tests	fix broken test	21 days ago	Releases 2			
	volumes	add link to download CTs	5 months ago	◊ v0.2 as used in new dataset p Latest			
	🗅 .gitignore	clean up scripts	4 months ago	on Jun 23			
	C README.md	Updated references for General Blockchain, Ir	nc. 9 days ago	+ 1 release			
	SCHEMA.md	Update findings table	7 days ago				
	🗅 metadata.csv	More RT-PCR annotations	2 days ago	Contributors 10			
				🐮 💷 📄 🙆 🕀 🍛 🚭			

Figure 4: Github Page to be able to download Data

# 5 Installing Required Library Modules

These are the python libraries that need to be pip installed and imported for the project.



Figure 5: Python Libraries & versions

# 6 Data Pre-processing

This is a part of pre-processing where the data is taken and has been divided into Test and Train sets.

s PC > Downloads > corona_dataset > Data								
Name	Date modified	Туре	Size					
📕 test	23-07-2020 10:02	File folder						
📕 train	23-07-2020 10:02	File folder						

#### Figure 6: Data divided into Training and Testing set

The Data is further divided into folders of Covid chest X-rays and Normal chest X-rays.

s PC > Downloads > corona_dataset > Data > test									
Name	Date modified	Туре	Size						
📙 Covid	23-07-2020 10:02	File folder							
📕 Normal	23-07-2020 10:02	File folder							

Figure 7: Data Divided into Covid & Normal



Figure 8: Covid Chest X-rays

### 7 Creating the Convolutional Neural Network (CNN)

Using tensorflow and Keras for CNN.

```
# Initialising the
 7 classifier = Sequential()
9 classifier.add(Conv2D(filters=32, kernel_size=3, padding="same", input_shape=(224, 224, 3),
                                                activation='relu'))
12 classifier.add(MaxPooling2D(pool_size=2, strides=2, padding='valid'))
14 classifier.add(Conv2D(filters=64, kernel_size=3, padding="same", activation="relu"))
15 classifier.add(MaxPooling2D(pool_size=2, strides=2, padding='valid'))
18 classifier.add(Conv2D(filters=64, kernel_size=3, padding="same", activation="relu"))
19 classifier.add(Dropout(0.2))
20 classifier.add(MaxPooling2D(pool_size=2, strides=2, padding='valid'))
23 classifier.add(Conv2D(filters=128, kernel_size=3, padding="same", activation="relu"))
24 classifier.add(Dropout(0.2))
25 classifier.add(MaxPooling2D(pool_size=2, strides=2, padding='valid'))
27 classifier.add(Conv2D(filters=256, kernel_size=3, padding="same", activation="relu"))
28 classifier.add(Dropout(0.2))
29 classifier.add(MaxPooling2D(pool_size=2, strides=2, padding='valid'))
31 classifier.add(Conv2D(filters=64, kernel_size=3, padding="same", activation="relu"))
32 classifier.add(MaxPooling2D(pool_size=2, strides=2, padding='valid'))
35 classifier.add(Flatten())
37 classifier.add(Dense(units=128, activation='relu'))
38 classifier.add(Dense(units=1, activation='sigmoid'))
40 classifier.compile(optimizer='adam', loss='binary_crossentropy', metrics=['accuracy'])
41 classifier.summary()
```

**Figure 9: CNN Model** 

# 8 Pre-processing and Loading Image set



Figure 10: Preprocessing and loading images

### **9** Training the Model





### 10 Passing the Test set through the model



#### Figure 12: Putting the test set through the model



Figure 13: Classifying the test set

# 11 Plotting Accuracy and Loss

```
In [21]: 1 #plot accuracy and Loss
2 import matplotlib.pyplot as plt
3
4 plt.plot(history.history['accuracy'])
5 plt.plot(history.history['val_accuracy'])
6 plt.title('model accuracy')
7 plt.ylabel('accuracy')
8 plt.xlabel('epoch')
9 plt.legend(['train', 'test'], loc='upper left')
10 plt.show()
11
12 plt.plot(history.history['loss'])
13 plt.plot(history.history['val_loss'])
14 plt.title('model loss')
15 plt.ylabel('loss')
16 plt.xlabel('epoch')
17 plt.legend(['train', 'test'], loc='upper left')
18 plt.show()
```

#### Figure 14: Plotting Accuracy & Loss of the Model

### 12 Classifying single images

```
2 import numpy as np
 3 from keras.preprocessing import image
5 test_image = image.load_img('../corona dataset/Data/test/Normal/IM-0283-0001.jpeg',target_size=(224,224
6 test_image = image.img_to_array(test_image)
 7 test_image = np.expand_dims(test_image, axis=0)
8 result = classifier.predict(test_image)
 9 training_set.class_indices
10 if result[0][0] == 1:
     prediction = 'Covid'
14 print(prediction)
Normal
 2 import numpy as np
 3 from keras.preprocessing import image
5 test_image = image.load_img('../corona dataset/Data/train/Covid/16654_1_1.png',target_size=(224,224))
6 test_image = image.img_to_array(test_image)
 7 test_image = np.expand_dims(test_image, axis=0)
8 result = classifier.predict(test_image)
 9 training_set.class_indices
10 if result[0][0] == 1:
       prediction = 'Normal'
       prediction = 'Covid'
14 print(prediction)
Covid
```

### **13** Plotting Confusion Matrix



**Figure 16: Plotting a Confusion Matrix** 

### 14 Evaluating Classification Report

The Classification Report[5] are necessary for Evaluation of any model.

In [26]: 1 from sklearn.metrics import classification_report									
2 3 print(c	lassifica	ation_rep	port(y_p	red, y_te	st))				
	precision	recall	f1-score	support					
	1.00	0.92	0.96	50					
	0.93	1.00	0.96	50					
accuracy			0.96	100					
macro avg	0.96	0.96	0.96	100					
weighted avg	0.96	0.96	0.96	100					

Figure 17: Classification Report

### References

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Cohen, J. P. (2020) *ieee8023/covid-chestxray-dataset*. Available at: https://github.com/ieee8023/covid-chestxray-dataset (Accessed: 15 August 2020).

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