1

Topic:-

Covid-19 Chest XRay Image Recognition

Problem Statement:-

Build an image classification model which will Identify covid poisitve patient by looking into the chest Xray images.

General Lifecycle of Image classification project

- 1) Data Collection
- 2) Importing libraries
- 3) Data Preprocessing
- 4) Model Building
- 5) Optimizing Model
- 6) Checking/ Vlidating model
- 7) Result

Importing all the importanat Library

In [1]:

```
import matplotlib.pyplot as plt
import cv2
import pathlib
from keras.layers import Dense, Conv2D, MaxPool2D, Dropout, Flatten
from keras.models import Sequential
```

In [28]:

```
#import imp lib
import numpy as np
import os
import pandas as pd
import tensorflow as tf
from tensorflow import keras
from tensorflow.keras import layers
import keras_tuner
from keras_tuner import RandomSearch
from keras_tuner.engine.hyperparameters import HyperParameters
from sklearn.metrics import f1_score,accuracy_score
from tensorflow.keras.metrics import Accuracy
from tensorflow.keras.preprocessing import image
from tensorflow.keras.preprocessing.image import ImageDataGenerator
```

Importing Data

About Dataset

Context Helping Deep Learning and Al Enthusiasts like me to contribute to improving COVID-19 detection using just Chest X-rays.

Content It is a simple directory structure branched into test and train and further branched into the respective 3 classes which contains the images.

Acknowledgements The University of Montreal for releasing the images.

Inspiration Help the medical and researcher community by sharing my work and encourage them to contribute extensively.

In [3]:

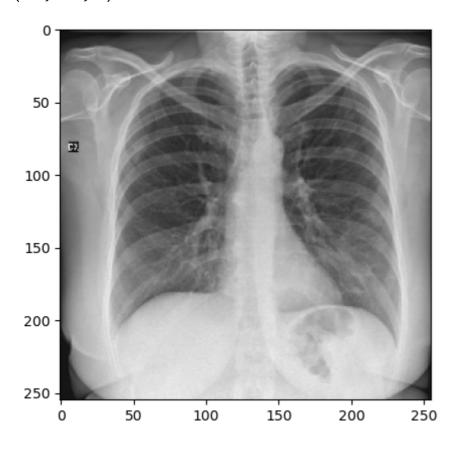
```
img = cv2.imread('C:/Users/Shivam Ranshur/ANACONDA JUPYTER PROGRAMMING/Covid19-datas
img = cv2.resize(img, (255, 255))
img = img / 255.0
```

In [4]:

```
plt.imshow(img)
img.shape
```

Out[4]:

(255, 255, 3)



Heatmap of xray just to show how covid Xray looks

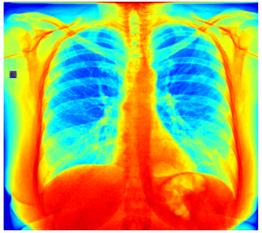
In [5]:

```
import cv2
   import numpy as np
   import matplotlib.pyplot as plt
 5
   # Load the image
 6 | image_path = 'C:/Users/Shivam Ranshur/ANACONDA JUPYTER PROGRAMMING/Covid19-dataset/t
 7
   image = cv2.imread(image_path)
9
   # Apply the colormap to the image
10
   heatmap = cv2.applyColorMap(image, cv2.COLORMAP_JET)
11
   # Convert the heatmap to RGB format
12
   heatmap_rgb = cv2.cvtColor(heatmap, cv2.COLOR_BGR2RGB)
13
14
15 # Display the original image and the heatmap overlay
16 fig, axs = plt.subplots(1, 2, figsize=(10, 5))
17
   axs[0].imshow(cv2.cvtColor(image, cv2.COLOR_BGR2RGB))
   axs[0].set_title('Original Image')
18
19
   axs[0].axis('off')
20 axs[1].imshow(heatmap_rgb)
   axs[1].set_title('Heatmap Overlay')
21
   axs[1].axis('off')
22
23
24
   plt.show()
25
```

Original Image



Heatmap Overlay



In [6]:

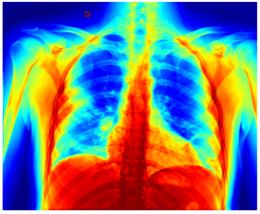
```
# Load the image
   image_path = "C:/Users/Shivam Ranshur/ANACONDA JUPYTER PROGRAMMING/Covid19-dataset/t
   image = cv2.imread(image_path)
 5
   # Apply the colormap to the image
   heatmap = cv2.applyColorMap(image, cv2.COLORMAP_JET)
 7
   # Convert the heatmap to RGB format
 9
   heatmap_rgb = cv2.cvtColor(heatmap, cv2.COLOR_BGR2RGB)
10
   # Display the original image and the heatmap overlay
11
12 fig, axs = plt.subplots(1, 2, figsize=(10, 5))
   axs[0].imshow(cv2.cvtColor(image, cv2.COLOR_BGR2RGB))
13
   axs[0].set_title('Original Image')
15 axs[0].axis('off')
16 | axs[1].imshow(heatmap_rgb)
   axs[1].set_title('Heatmap Overlay')
17
   axs[1].axis('off')
18
19
20
   plt.show()
21
```



Original Image



Heatmap Overlay



Data PreProcessing

In [7]:

1 | # we have 251 images in training data which is very less for CNN so we need to do da 2 | # Data augmentation is a technique to create more data using images we have currentl 3 # we'll do this on training data

Data Augmentation

In [9]:

```
1
   # Data augmention is use to generate more data using exixting data
   train_data = "C:/Users/Shivam Ranshur/ANACONDA JUPYTER PROGRAMMING/Covid19-dataset/t
   target_size = (256, 256)
   batch_size = 16
 5
 6
   train_datagen = ImageDataGenerator(
 7
        rescale=1 / 255,
 8
        rotation_range=0.2,
 9
        width shift range=0.1,
10
        height_shift_range=0.1,
11
        shear_range=0.2,
        zoom_range=0.2,
12
13
        horizontal_flip=True,
        fill_mode='nearest'
14
15
    )
```

In [10]:

```
1
  train_generator = train_datagen.flow_from_directory(
2
      train_data,
3
      target_size=target_size,
4
      batch_size=batch_size,
5
      class_mode='categorical'
6
  )
7
  # Calculate the total number of samples in the generator
  num_samples = len(train_generator) * batch_size
9
  print("Total number of samples in the ImageDataGenerator:", num_samples)
```

Found 251 images belonging to 3 classes.

Total number of samples in the ImageDataGenerator: 256

Getting our labels

In [11]:

```
# other way is
train_generator.class_indices # we got our labels of data
```

```
Out[11]:
```

```
{'Covid': 0, 'Normal': 1, 'Viral Pneumonia': 2}
```

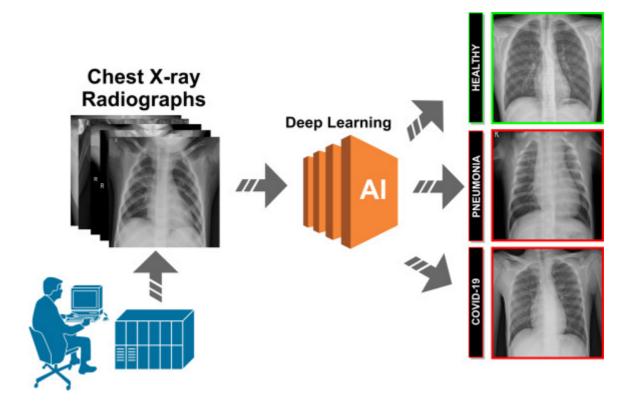
```
In [12]:
```

```
# doing same for test data
    # we could have keep same data augmentation code for test and train both , but for e
    test_data = "C:/Users/Shivam Ranshur/ANACONDA JUPYTER PROGRAMMING/Covid19-dataset/te
    target_size = (256, 256)
 5
    batch_size = 16
 6
 7
    test_datagen = ImageDataGenerator(
 8
        rescale=1 / 255,
 9
        rotation_range=0.2,
10
        width shift range=0.1,
        height_shift_range=0.1,
11
12
        shear_range=0.2,
13
        zoom_range=0.2,
14
        horizontal_flip=True,
        fill_mode='nearest'
15
16
    )
17
18
    test_generator = test_datagen.flow_from_directory(
19
20
        test_data,
21
        target_size=target_size,
22
        batch_size=batch_size,
23
        class_mode='categorical'
24
    )
25
26 # Calculate the total number of samples in the generator
    num_samples = len(test_generator) * batch_size
    print("Total number of samples in the ImageDataGenerator:", num_samples)
Found 66 images belonging to 3 classes.
Total number of samples in the ImageDataGenerator: 80
In [13]:
    test_generator.class_indices # labels for test data
Out[13]:
{'Covid': 0, 'Normal': 1, 'Viral Pneumonia': 2}
In [ ]:
 1
```

MODEL(CNN)

In [14]:

```
Xray model = Sequential()
 2
   Xray_model.add(Conv2D(filters=32, kernel_size=(3,3), activation="relu",input_shape=(
4
   Xray model.add(MaxPool2D())
   Xray_model.add(Dropout(rate=0.23))#Dropout is use to prevent overfitting
 5
 6
 7
   Xray_model.add(Conv2D(filters=64, kernel_size=(3,3), activation="relu"))
   Xray_model.add(MaxPool2D())
8
9
   Xray_model.add(Dropout(rate=0.25))
10
   Xray_model.add(Conv2D(filters=128, kernel_size=(3,3), activation="relu"))
11
   Xray model.add(MaxPool2D())
12
   Xray_model.add(Dropout(rate=0.25))
13
14
15  Xray_model.add(Flatten())
16 Xray_model.add(Dense(units = 64, activation="relu"))
   Xray_model.add(Dropout(rate=0.40))
17
   Xray_model.add(Dense(units=3, activation="softmax"))# 3 inits is used because we hav
18
19
   Xray_model.compile(loss="categorical_crossentropy", optimizer= "adam",metrics= ["acc
20
```



Model summary

In [15]:

1 Xray_model.summary()

Model: "sequential"

model: Sequential		
Layer (type)	Output Shape	Param #
conv2d (Conv2D)	(None, 254, 254, 32)	896
<pre>max_pooling2d (MaxPooling2D)</pre>	(None, 127, 127, 32)	0
dropout (Dropout)	(None, 127, 127, 32)	0
conv2d_1 (Conv2D)	(None, 125, 125, 64)	18496
<pre>max_pooling2d_1 (MaxPooling 2D)</pre>	(None, 62, 62, 64)	0
dropout_1 (Dropout)	(None, 62, 62, 64)	0
conv2d_2 (Conv2D)	(None, 60, 60, 128)	73856

Model fitting

In [21]:

1 Xray_model.fit(train_generator,epochs=20,steps_per_epoch= 8,validation_steps=2)

```
Epoch 1/20
cy: 0.8594
Epoch 2/20
8/8 [=========== ] - 16s 2s/step - loss: 0.3279 - accura
cy: 0.8750
Epoch 3/20
cy: 0.8750
Epoch 4/20
cy: 0.8906
Epoch 5/20
cy: 0.8537
Epoch 6/20
cy: 0.8984
Epoch 7/20
cy: 0.9106
Epoch 8/20
cy: 0.8780
Epoch 9/20
cy: 0.8906
Epoch 10/20
cy: 0.9024
Epoch 11/20
cy: 0.9106
Epoch 12/20
cy: 0.8374
Epoch 13/20
cy: 0.8699
Epoch 14/20
cy: 0.8984
Epoch 15/20
cy: 0.8594
Epoch 16/20
cy: 0.8438
Epoch 17/20
cy: 0.8862
Epoch 18/20
cy: 0.9268
Epoch 19/20
cy: 0.9024
Epoch 20/20
cy: 0.9106
```

Out[21]:

<keras.callbacks.History at 0x185494ec130>

```
In [ ]:
```

```
# our model is giving final accuracy of 91.06
#which is fine
#we can optimize model further but it is not always necessary to complicate models
# if it is working perfectly consuming less computational power
# if while validating it dosent poerform well well optimize it further
```

In [22]:

```
1 y_pred = Xray_model.predict(test_generator)
```

```
5/5 [======] - 5s 864ms/step
```

Validating Model

We'll do this by passing images of different categories

Labels

{'Covid': 0, 'Normal': 1, 'Viral Pneumonia': 2}

In [30]:

```
test_path= "C:/Users/Shivam Ranshur/ANACONDA JUPYTER PROGRAMMING/Covid19-dataset/tra
img= image.load_img(test_path, target_size=(256,256))
img = image.img_to_array(img)/255
img= np.array([img])
print(img.shape)
```

(1, 256, 256, 3)

In [32]:

```
# Predict the class probabilities for the image
class_probabilities = Xray_model.predict(img)

# Get the predicted class label
predicted_class = np.argmax(class_probabilities, axis=1)[0]

print("Class Probabilities:", class_probabilities)
print("Predicted Class:", predicted_class)
```

```
1/1 [============== ] - 0s 276ms/step Class Probabilities: [[0.98729765 0.01164208 0.00106034]] Predicted Class: 0
```

```
In [34]:
```

```
# we can see name in path its normal
 2 test_path= "C:/Users/Shivam Ranshur/ANACONDA JUPYTER PROGRAMMING/Covid19-dataset/tes
 3 img= image.load_img(test_path, target_size=(256,256))
 4 img = image.img_to_array(img)/255
   img= np.array([img])
 6 print(img.shape)
 7
 8 # Predict the class probabilities for the image
 9
   class_probabilities = Xray_model.predict(img)
10
11
   # Get the predicted class label
12
   predicted_class = np.argmax(class_probabilities, axis=1)[0]
13
14 | print("Class Probabilities:", class_probabilities)
   print("Predicted Class:", predicted_class)
15
(1, 256, 256, 3)
1/1 [======= ] - 0s 28ms/step
Class Probabilities: [[4.7880650e-04 6.5907425e-01 3.4044698e-01]]
Predicted Class: 1
In [35]:
 1 | # we can see name in path its pneunomia
   test_path= "C:/Users/Shivam Ranshur/ANACONDA JUPYTER PROGRAMMING/Covid19-dataset/tes
   img= image.load_img(test_path, target_size=(256,256))
 4 | img = image.img to array(img)/255
 5 img= np.array([img])
   print(img.shape)
 7
   # Predict the class probabilities for the image
 9
   class_probabilities = Xray_model.predict(img)
10
11 # Get the predicted class label
predicted_class = np.argmax(class_probabilities, axis=1)[0]
13
14
   print("Class Probabilities:", class_probabilities)
```

print("Predicted Class:", predicted_class)

For all 3 categories its working perfectely as we wanted its classifying perfectly

RESULT

Proposed CNN model achieves 91% accuracy in X-ray image classification, effectively detecting COVID-19, normal, and pneumonia cases with high precision.

Feel free to use this project and tune it further for your personal projects