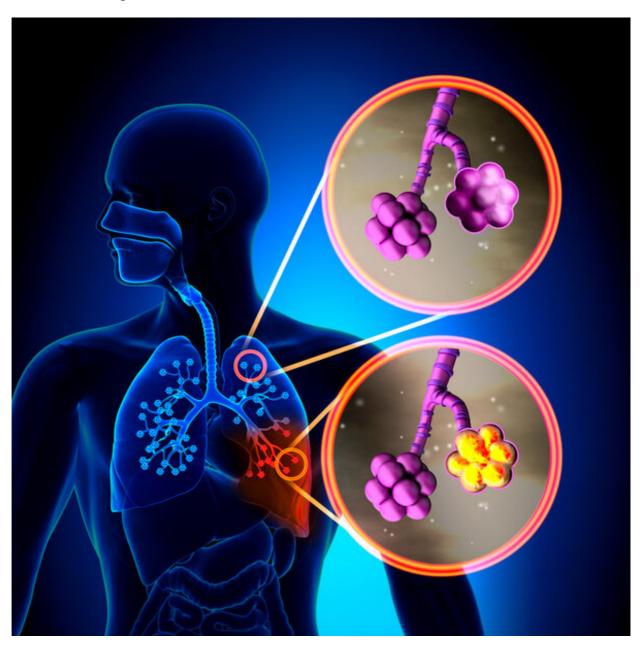
Author: Namita Rana

## **Project Name**

Detection of Pneumonia from Chest X-Ray Images using ConvolutionalNeural Network and Transfer Learning.



#### **Dataset Details**

Dataset Name: Chest X-Ray Images (Pneumonia)

Number of Class: 2

Number/Size of Images: Total: 5863 Training: 5216 Validation: 16

## **Business Problem:**

Building a model that can classify whether a given patient has pneumonia, given a chest x-ray image.

Stakeholer: Imaging labs/ Hospitals.

Business Questions: How can a successful model help save medical professionals time, money and promote better accuracy in patient diagnosis.

## **Background:**

Pneumonia is an infection that inflames your lungs' air sacs (alveoli). The air sacs may fill up with fluid or pus, causing symptoms such as a cough, fever, chills and trouble breathing. Bacteria and viruses are the main causes of pneumonia. Pneumonia-causing germs can settle in the alveoli and multiply after a person breathes them in. Pneumonia can be contagious. The bacteria and viruses that cause pneumonia are usually inhaled. Commonly affected are Infants, children and people over 65 years in age.

Chest X-rays are used for detecting the Pneumonia infection and to locate the infected area in the lungs. So, To detect the the pneumonia radiologist have to observe the chest xray and he/she has to update the doctor correctly. The main objective of this model is to identify if the person has Pneumonia or not with high accuracy so that the person can get treatment as soon as possible. Deep Learning models which are trained correctly by using good datasets can be helpful for doctors.

To train the model for detecting whether the person has pneumonia or not, A Convolutional Neural Network(CNN) is used. The CNN can train the images of chest xrays and then it can predict with high accuracy.

## **Data Structure, Selection & Transformation:**

The dataset we recieved in Kaggle is actually distributed into 3 folders (train, test, val) and individually, they contain subfolders for each image category (Pneumonia/Normal).

There are a total of 5,863 X-Ray images (in JPEG Format) distributed into 2 categories (Pneumonia/Normal).

Chest X-ray images (anterior-posterior) were selected from retrospective cohorts of pediatric patients of one to five years old from Guangzhou Women and Children's Medical Center, Guangzhou. All chest X-ray imaging was performed as part of patients' routine clinical care. For the analysis of chest x-ray images, all chest radiographs were initially screened for quality control by removing all low quality or unreadable scans. The diagnoses for the images were then graded by two expert physicians before being cleared for training the Al system. In order to account for any grading errors, the evaluation set was also checked by a third expert.

### **Methods**

# **Cleaning and Feature Engineering**

This project uses data cleaning and feature engineering to also addressed the class imbalance between classes we have used Data Augmentation.

Data augmentation in data analysis are techniques used to increase the amount of data by adding slightly modified copies of already existing data or newly created synthetic data from existing data. It acts as a regularizer and helps reduce overfitting when training a machine learning model.

In order to avoid overfitting problem, we need to expand artificially our dataset. We can make your existing dataset even larger. The idea is to alter the training data with small transformations to reproduce the variations. Approaches that alter the training data in ways that change the array representation while keeping the label the same are known as data augmentation techniques. Some popular augmentations people use are grayscales, horizontal flips, vertical flips, random crops, color jitters, translations, rotations, and much more. By applying just a couple of these transformations to our training data, we can easily double or triple the number of training examples and create a very robust model.

## **Models Development**

We have implemented versions of CNN's with different parameters, dense layers, dropout layers to see how results varies with each change in the parameters. Results Our model, CNN came back with a confusion matrix that produced a 91% accuracy score and a 99% recall score. For our purposes, we were looking to minimize recall as we want to reduce the amount of False positives (False negatives: Patients got negative results but has actually has Pneumonia).

#### **Metrics used:**

- 1. Accuracy
- 2. Recall

Here, "Recall" is the most significant metric even more than accuracy and precision. False negative has to be minimized because falsely diagnosing a patient of pneumonia as not having pneumonia is a much larger concern than falsely diagnosing a healthy person as a pneumonia patient. By minimizing false negative, which is in the denominator, we can increase 'Recall' .This model achieves a Recall of 99%.

## **Import Packages and Functions**

We'll make use of the following packages:

numpy and pandas is what we'll use to manipulate our data.

matplotlib.pyplot and seaborn will be used to produce plots for visualization.

util will provide the locally defined utility functions that have been provided for this assignment.

We will also use several modules from the keras framework for building deep learning models.

Run the next cell to import all the necessary packages.

```
In [3]:
            #Importing all the necessary libraries:
         1
            #we will be using keras for the Neural net stuff.
           import os
            import pandas as pd
           import numpy as np
           import matplotlib.pyplot as plt
            import seaborn as sns
           import keras
            from keras.models import Sequential
        10 from keras.layers import Dense, Conv2D , MaxPool2D , Flatten , Drd
           from keras.preprocessing.image import ImageDataGenerator
           from sklearn.model selection import train test split
        12
           from sklearn.metrics import classification report, confusion matrix
        14
           from keras.callbacks import ReduceLROnPlateau
        15 import cv2
            import datetime
```

## Reading the data images

Creating a helper function to pull in the data. Using CV2 for this. We will pull in the images in a 150\*150.

```
In [4]:
            #helper function:
            labels = ['PNEUMONIA', 'NORMAL']
          2
            img size = 150
            def get data(data dir):
          5
                data = []
                for label in labels:
          6
          7
                     path = os.path.join(data dir, label)
                     class_num = labels.index(label)
          8
                     for img in os.listdir(path):
          9
        10
                         try:
         11
                             img arr = cv2.imread(os.path.join(path, img), cv2.
                             resized arr = cv2.resize(img arr, (img size, img s
        12
                             data.append([resized arr, class num])
        13
         14
                         except Exception as e:
        15
                             print(e)
                return np.array(data)
         16
```

```
In [5]: 1 os.listdir("chest_xray")
Out[5]: ['.DS_Store', 'test', 'chest_xray', '__MACOSX', 'train', 'val']
In [6]: 1 len(os.listdir("chest_xray/train/PNEUMONIA"))
Out[6]: 3875
```

## Loading the dataset.

```
#Using get_data function to load the dataset.
In [7]:
           train = get data('chest xray/chest xray/train')
        2
          test = get data('chest xray/chest xray/test')
        3
           val = get data('chest xray/chest xray/val')
           print("Train set:\n========")
        7
          num pneumonia = len(os.listdir("chest xray/train/PNEUMONIA"))
        8 num normal = len(os.listdir("chest xray/train/NORMAL"))
           print(f"PNEUMONIA={num pneumonia}")
          print(f"NORMAL={num normal}")
       10
       11
       12
          print("Test set:\n=========")
           num pneumonia = len(os.listdir("chest xray/test/PNEUMONIA"))
          num normal = len(os.listdir("chest xray/test/NORMAL"))
       14
       15
           print(f"PNEUMONIA={num pneumonia}")
          print(f"NORMAL={num normal}")
       16
       17
       18
           print("Val set:\n==========")
          num pneumonia = len(os.listdir("chest xray/val/PNEUMONIA"))
       20
           num normal = len(os.listdir("chest xray/val/NORMAL"))
           print(f"PNEUMONIA={num pneumonia}")
       21
       22
           print(f"NORMAL={num normal}")
       23
```

OpenCV(4.5.5) /private/var/folders/\_x/m61x8cs124q78tmfkx9svwzw0000gn /T/pip-install-fecr2822/opencv-python/opencv/modules/imgproc/src/res ize.cpp:4052: error: (-215:Assertion failed) !ssize.empty() in funct ion 'resize'

OpenCV(4.5.5) /private/var/folders/\_x/m61x8cs124q78tmfkx9svwzw0000gn /T/pip-install-fecr2822/opencv-python/opencv/modules/imgproc/src/res ize.cpp:4052: error: (-215:Assertion failed) !ssize.empty() in funct ion 'resize'

OpenCV(4.5.5) /private/var/folders/\_x/m61x8cs124q78tmfkx9svwzw0000gn /T/pip-install-fecr2822/opencv-python/opencv/modules/imgproc/src/res ize.cpp:4052: error: (-215:Assertion failed) !ssize.empty() in funct ion 'resize'

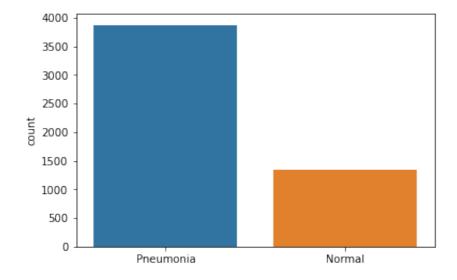
OpenCV(4.5.5) /private/var/folders/\_x/m6lx8cs124q78tmfkx9svwzw0000gn

```
/T/pip-install-fecr2822/opencv-python/opencv/modules/imgproc/src/res
       ize.cpp:4052: error: (-215:Assertion failed) !ssize.empty() in funct
       ion 'resize'
       Train set:
       PNEUMONIA=3875
      NORMAL=1341
      Test set:
       PNEUMONIA=390
      NORMAL=234
      Val set:
       _____
       PNEUMONIA=8
      NORMAL=8
          pneumonia = os.listdir("chest_xray/train/PNEUMONIA")
In [8]:
         pneumonia dir = "chest xray/train/PNEUMONIA"
```

Let us try to see how well is the data distributed.

/Users/Ravinder/opt/anaconda3/envs/learn-env/lib/python3.8/site-pack ages/seaborn/\_decorators.py:36: FutureWarning: Pass the following variable as a keyword arg: x. From version 0.12, the only valid positi onal argument will be `data`, and passing other arguments without an explicit keyword will result in an error or misinterpretation. warnings.warn(

Out[9]: <AxesSubplot:ylabel='count'>



We can see that we have an imbalanced dataset. Hence, we will be using Data Augmentation to increase the no of training samples.

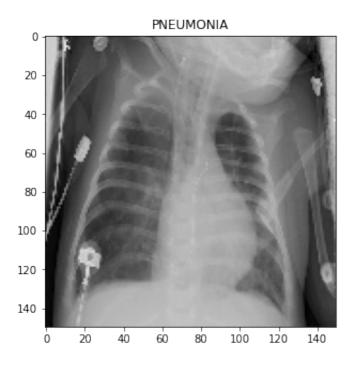
# Previewing the images of both the classes.

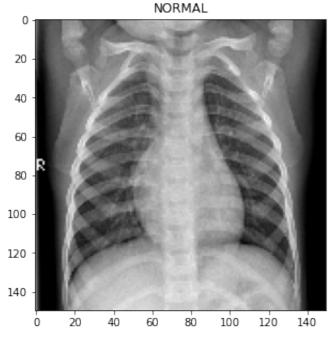
```
In [10]: #plot to view images of both Pnumeonia & Normal class.

2 plt.figure(figsize = (5,5))
    plt.imshow(train[0][0], cmap='gray')
    plt.title(labels[train[0][1]])

5 plt.figure(figsize = (5,5))
    plt.imshow(train[-1][0], cmap='gray')
    plt.title(labels[train[-1][1]])
```

Out[10]: Text(0.5, 1.0, 'NORMAL')





### **Data Preparation:**

• In image classification we pull in images into an array of numbers. These numbers represent the pixel intensity, and it's a number between 0 and 255, 255 being white and 0 being black.

#### Points to be considered:

- We need to pair the arrays of the image with their labels (if pneumonia or not)
- In a neural net, there's a lot of math that happens in the backend. If we use numbers like 255 the computer is forced to work with really huge numbers which may increase computational power as well as slow down the epochs. To rectify this we can just divide every pixel by 255 that way we're left with numbers between 0 and 1, 1 being white and 0 being black.
- Lastly when we feed the images to keras, we need to reshape our dimensions. So we'll use the x\_train.reshape(-1, image\_size, image\_size, 1). The numbers mean [batch\_size, height, width, channels]. The -1 means that the length in the dimension is inferred so we don't have to specify it. The 1 is because we're using a black and white picture so we'll only have one layer image.

```
In [12]:
             #Pairing the array of the image with their labels(if pneumonia or
           2
             x train = []
           3
             y train = []
           4
           5
             for feature, label in train:
           6
                  x train.append(feature)
           7
                  y_train.append(label)
           8
             #print(x train)
           9
             x val = []
          10
             y val = []
          11
          12
             for feature, label in val:
                  x val.append(feature)
          13
          14
                  y val.append(label)
          15
          16
             x test = []
          17
             y test = []
          18
          19
             for feature, label in test:
          20
                  x test.append(feature)
          21
                  y test.append(label)
```

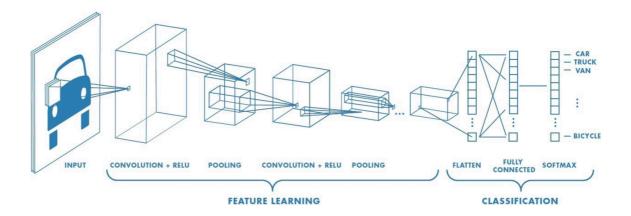
```
# Normalize the data
In [13]:
           1
           2
              x train = np.array(x train) / 255
             x_val = np.array(x_val) / 255
           3
              x \text{ test} = \text{np.array}(x \text{ test}) / 255
In [14]:
              #Resizing the data for deep learning
           1
           2
              x_train = x_train.reshape(-1, img_size, img size, 1)
           3
             y_train = np.array(y_train)
           4
             x_val = x_val.reshape(-1, img_size, img_size, 1)
             y val = np.array(y val)
           8
             x test = x test.reshape(-1, img size, img size, 1)
              y_test = np.array(y_test)
              print(x train[0][0].shape)
```

### What is CNN?

(150, 1)

CNN stands for Convolutional Neural Network which is a specialized neural network for processing data that has an input shape like a 2D matrix like images. CNN's are typically used for image detection and classification.

The architecture of a ConvNet is analogous to that of the connectivity pattern of Neurons in the Human Brain and was inspired by the organization of the Visual Cortex. Individual neurons respond to stimuli only in a restricted region of the visual field known as the Receptive Field. A collection of such fields overlap to cover the entire visual area.



```
In [15]: # Build a baseline fully connected model
2  from keras import models
3  from keras import layers
4  #np.random.seed(123)
5  model = models.Sequential()
6  model.add(layers.Dense(20, activation='relu', input_shape=(150,150, model.add(layers.Dense(7, activation='relu'))
8  model.add(layers.Dense(5, activation='relu'))
9  model.add(layers.Dense(1, activation='sigmoid'))
```

### Fixing data imbalance: Data Augmentation

- We will create duplicates of the images we have by doing certain changes to the images, which can include rotating, flipping, zooming, shifting etc.
- We will not be performing flipping as all the x-Rays will be vertical so doing that might be confusing for the computer.

```
In [17]: m the data augmentation, here the ImageDataGenerator class provided by
        ImageDataGenerator(
        aturewise center=False,
         mplewise center=False,
         aturewise std normalization=False,
        mplewise std normalization=False,
         a whitening=False,
         tation range = 30,
         om 9range = 0.2,
        dth shift range=0.1,
        idht shift range=0.1,
        rizontal_flip = True,
        rtlBcal flip=False)
         t(1x train)
In [18]:
          1 #timing the model:
           2 original start = datetime.datetime.now()
             start = datetime.datetime.now()
```

### CNN1 with a dropout layer.

```
In [19]: nstantiating the model

del = Sequential()

dding the first layer

tep1: Convolution

del. add(Conv2D(32,(3,3),strides =1,padding='same',activation ='relu',ir

tep2: Pooling

del. add(MaxPool2D((2,2)))

8

dding a second convolutional layer
```

```
deD.add(Conv2D(64,(3,3),strides =1,padding='same',activation ='relu'))
dell.add(Dropout(0.1))
oolinh
d \in \mathbb{B}. add(MaxPool2D((2,2)))
 14
ddfing a third convolutional layer
deB.add(Conv2D(128,(3,3),strides =1,padding='same',activation ='relu')
deT.add(Dropout(0.3))
odBing
d \in \mathbb{P}. add(MaxPool2D((2,2)))
daing a fourth layer
del.add(Conv2D(128,(3,3),strides =1,padding='same',activation ='relu')
deB.add(Dropout(0.2))
d@1.add(MaxPool2D((2,2)))
 25
ddbng a fifth layer
deT.add(Conv2D(256,(3,3),strides =1,padding='same',activation ='relu')
deB.add(Dropout(0.2))
d@\mathbb{P}. add(MaxPool2D((2,2)))
tep3: Flattening
dell.add(Flatten())
deB.add(Dense(units =128,activation = 'relu'))
dell.add(Dropout(0.3))
 35
tep4: Full connection
deT.add(Dense(units=1,activation ='sigmoid'))
ombling the CNN
deD.compile(optimizer = "adam",loss = binary crossentropy',metrics = ['d
isplay model summary.
del.summary()
```

Model: "sequential\_1"

Layer (type)	Output	Shape	Param #
conv2d (Conv2D)	(None,	150, 150, 32)	320
max_pooling2d (MaxPooling2D)	(None,	75, 75, 32)	0
conv2d_1 (Conv2D)	(None,	75, 75, 64)	18496
dropout (Dropout)	(None,	75, 75, 64)	0
max_pooling2d_1 (MaxPooling2	(None,	37, 37, 64)	0
conv2d_2 (Conv2D)	(None,	37, 37, 128)	73856
dropout_1 (Dropout)	(None,	37, 37, 128)	0
<pre>max_pooling2d_2 (MaxPooling2</pre>	(None,	18, 18, 128)	0

conv2d_3 (Conv2D)	(None,	18, 18, 128)	147584
dropout_2 (Dropout)	(None,	18, 18, 128)	0
max_pooling2d_3 (MaxPooling2	(None,	9, 9, 128)	0
conv2d_4 (Conv2D)	(None,	9, 9, 256)	295168
dropout_3 (Dropout)	(None,	9, 9, 256)	0
max_pooling2d_4 (MaxPooling2	(None,	4, 4, 256)	0
flatten (Flatten)	(None,	4096)	0
dense_4 (Dense)	(None,	128)	524416
dropout_4 (Dropout)	(None,	128)	0
dense_5 (Dense)	(None,	1)	129

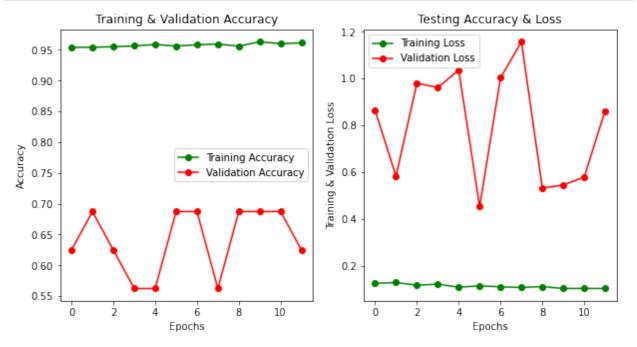
Total params: 1,059,969
Trainable params: 1,059,969
Non-trainable params: 0

```
In [64]: #Fitting the CNN ro the images using fit_generator
history = model.fit(datagen.flow(x_train,y_train,batch_size=32),epoc

# end timer
end = datetime.datetime.now()
elapsed = end - start
print('Training took a total of {}'.format(elapsed))
```

```
Epoch 1/12
41 - accuracy: 0.9542 - val loss: 0.8646 - val accuracy: 0.6250
75 - accuracy: 0.9544 - val loss: 0.5824 - val accuracy: 0.6875
Epoch 3/12
62 - accuracy: 0.9551 - val loss: 0.9804 - val accuracy: 0.6250
Epoch 4/12
07 - accuracy: 0.9565 - val loss: 0.9620 - val accuracy: 0.5625
Epoch 5/12
77 - accuracy: 0.9588 - val loss: 1.0364 - val accuracy: 0.5625
Epoch 6/12
34 - accuracy: 0.9563 - val loss: 0.4528 - val accuracy: 0.6875
Epoch 7/12
92 - accuracy: 0.9584 - val loss: 1.0031 - val accuracy: 0.6875
64 - accuracy: 0.9595 - val loss: 1.1574 - val accuracy: 0.5625
Epoch 9/12
03 - accuracy: 0.9561 - val loss: 0.5323 - val accuracy: 0.6875
Epoch 10/12
21 - accuracy: 0.9632 - val loss: 0.5444 - val accuracy: 0.6875
Epoch 11/12
24 - accuracy: 0.9601 - val loss: 0.5786 - val accuracy: 0.6875
Epoch 12/12
20 - accuracy: 0.9615 - val loss: 0.8618 - val accuracy: 0.6250
Training took a total of 5:31:46.546904
```

```
In [67]:
             #plotting the Training accuracy and loss
             epochs = [i for i in range(12)]
           2
             fig , ax = plt.subplots(1,2)
           3
             train_acc = history.history['accuracy']
             train loss = history.history['loss']
           5
             val acc = history.history['val accuracy']
           7
             val loss = history.history['val loss']
             fig.set size inches(10,5)
           8
          9
          10
             ax[0].plot(epochs , train_acc , 'go-' , label = 'Training Accuracy
             ax[0].plot(epochs , val_acc , 'ro-' , label = 'Validation Accuracy
          11
          12
             ax[0].set title('Training & Validation Accuracy')
         13
             ax[0].legend()
          14
             ax[0].set xlabel("Epochs")
             ax[0].set ylabel("Accuracy")
         15
          16
         17
             ax[1].plot(epochs , train_loss , 'g-o' , label = 'Training Loss')
             ax[1].plot(epochs , val loss , 'r-o' , label = 'Validation Loss')
          18
          19
             ax[1].set title('Testing Accuracy & Loss')
          20
             ax[1].legend()
          21
             ax[1].set xlabel("Epochs")
             ax[1].set ylabel("Training & Validation Loss")
          23
             plt.show()
          24
```



#### **Predictions**

```
In [69]:
              cm = confusion matrix(y test,predictions)
              cm = pd.DataFrame(cm , index = ['0', '1'] , columns = ['0', '1'])
In [70]: lt.lfigure(figsize = (5,5))
         ns 2heatmap(cm,cmap= "Blues", linecolor = 'black', linewidth = 1, anno
Out[70]: <AxesSubplot:>
                                                350
                                               300
                    385
                                   5
           PNEUMONIA
                                                250
                                                200
                                               - 150
                    53
                                  181
                                               - 100
           NORMAL
                                               - 50
                 PNEUMONIA
                                 NORMAL
In [71]:
              print(classification report(y test,predictions,target names = ['Pr
                                precision
                                               recall
                                                        f1-score
                                                                    support
          Pneumonia(class 0)
                                      0.88
                                                 0.99
                                                            0.93
                                                                         390
                                                 0.77
             Normal(Class 1)
                                      0.97
                                                            0.86
                                                                         234
                                                            0.91
                                                                         624
                     accuracy
                    macro avg
                                      0.93
                                                 0.88
                                                            0.90
                                                                         624
                 weighted avg
                                      0.91
                                                 0.91
                                                            0.90
                                                                         624
In [72]:
              #save the model
```

#### **Results:**

Model with dropout layers has an accuracy of %, and recall of 99% on Recall.

model.save("model.h5")

# Now's let build a CNN without drop out layer:

We want to see how our model behaves without dropout layers.

### CNN<sub>2</sub>

```
In [24]: Buildign the CNN2 model
        del2 = Sequential()
        Adding the first layer
        Stepl: Convolution
        bdell2.add(Conv2D(32,(3,3),strides =1,padding='same',activation ='relu',
        Step2:Pooling
         de12.add(MaxPool2D((2,2)))
         Adding a second convolutional layer
        bdel2.add(Conv2D(64,(3,3),strides =1,padding='same',activation ='relu')
         nodel.add(Dropout(0.2))
         Pob2L1.nh
         del2.add(MaxPool2D((2,2)))
         Adding a third convolutional layer
         bde12.add(Conv2D(64,(3,3),strides =1,padding='same',activation ='relu')
         nodel.add(Dropout(0.3))
         Pob&Ling
        del2.add(MaxPool2D((2,2)))
         Adding a fourth convolutional layer
         bd@12.add(Conv2D(128,(3,3),strides =1,padding='same',activation ='relu'
         nodel.add(Dropout(0.3))
         PooAing
        del2.add(MaxPool2D((2,2)))
          2.6
         Adding a fifth convolutional layer
         del2.add(Conv2D(128,(3,3),strides =1,padding='same',activation ='relu'
         nodel.add(Dropout(0.3))
         Pooling
         del2.add(MaxPool2D((2,2)))
          32
         adding a sixth layer
        bde42.add(Conv2D(128,(3,3),strides =1,padding='same',activation ='relu'
         nodel.add(Dropout(0.2))
        ded2.add(MaxPool2D((2,2)))
          37
         Step3: Flattening
        del2.add(Flatten())
        bde12.add(Dense(units =256,activation = 'relu'))
        dell2.add(Dropout(0.3))
         Step4: Full connection
        bde12.add(Dense(units=1,activation ='sigmoid'))
         Comming the CNN
```

```
inded2.compile(optimizer = "adam",loss = 'binary_crossentropy',metrics =[
iispi ay model summary.
)ded2.summary()
49
```

Model: "sequential\_2"

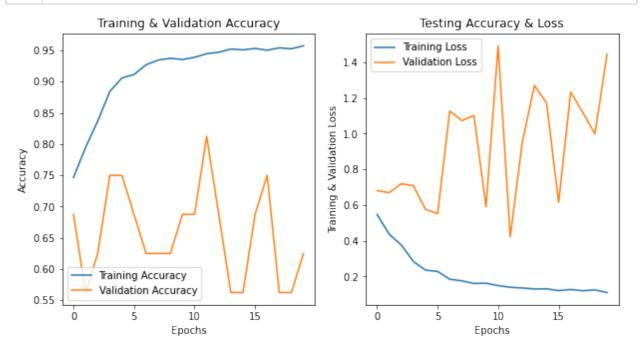
Output	Shape	Param #
(None,	150, 150, 32)	320
(None,	75, 75, 32)	0
(None,	75, 75, 32)	9248
(None,	37, 37, 32)	0
(None,	37, 37, 64)	18496
(None,	18, 18, 64)	0
(None,	18, 18, 64)	36928
(None,	9, 9, 64)	0
(None,	9, 9, 128)	73856
(None,	4, 4, 128)	0
(None,	4, 4, 128)	147584
(None,	2, 2, 128)	0
(None,	512)	0
(None,	256)	131328
(None,	256)	0
(None,	1)	257 =======
	(None,	Output Shape  (None, 150, 150, 32)  (None, 75, 75, 32)  (None, 37, 37, 32)  (None, 37, 37, 64)  (None, 18, 18, 64)  (None, 18, 18, 64)  (None, 9, 9, 64)  (None, 9, 9, 128)  (None, 4, 4, 128)  (None, 4, 4, 128)  (None, 2, 2, 128)  (None, 512)  (None, 256)  (None, 256)

Total params: 418,017 Trainable params: 418,017 Non-trainable params: 0

```
In [25]:
       1 #Fitting the CNN ro the images using fit generator
        2 history = model2.fit(datagen.flow(x_train,y_train,batch_size=32),e
       3
        4
         # end timer
         end = datetime.datetime.now()
         elapsed = end - start
         print('Training took a total of {}'.format(elapsed))
      Epoch 10/20
      0.1623 - accuracy: 0.9354 - val loss: 0.5906 - val accuracy: 0.687
      Epoch 11/20
      0.1490 - accuracy: 0.9390 - val loss: 1.4920 - val accuracy: 0.687
      Epoch 12/20
      0.1397 - accuracy: 0.9448 - val loss: 0.4222 - val accuracy: 0.812
      Epoch 13/20
      0.1356 - accuracy: 0.9471 - val loss: 0.9536 - val_accuracy: 0.687
      Epoch 14/20
      0.1302 - accuracy: 0.9523 - val loss: 1.2690 - val accuracy: 0.562
       1 print("Loss of the model is - " , model2.evaluate(x_test,y_test)[0]
In [27]:
        2 print("Accuracy of the model is - " , model2.evaluate(x test,y tes
      20/20 [=============== ] - 4s 206ms/step - loss: 0.340
      8 - accuracy: 0.9119
      Loss of the model is - 0.34082722663879395
      20/20 [=============== ] - 4s 212ms/step - loss: 0.340
      8 - accuracy: 0.9119
```

Accuracy of the model is - 91.18589758872986 %

```
#plotting training& validation accuracy, Traning & validation loss
In [28]:
             import matplotlib.pyplot as plt
           2
           3
             epochs = [i for i in range(20)]
             fig , ax = plt.subplots(1,2)
             train acc = history.history['accuracy']
             train loss = history.history['loss']
           7
             val acc = history.history['val accuracy']
             val loss = history.history['val loss']
          9
             fig.set size inches(10,5)
          10
             ax[0].plot(epochs , train_acc , label = 'Training Accuracy')
          11
             ax[0].plot(epochs , val_acc , label = 'Validation Accuracy')
         12
         13
             ax[0].set title('Training & Validation Accuracy')
          14
             ax[0].legend()
             ax[0].set xlabel("Epochs")
         15
             ax[0].set_ylabel("Accuracy")
          16
         17
             ax[1].plot(epochs , train loss , label = 'Training Loss')
         18
             ax[1].plot(epochs , val loss , label = 'Validation Loss')
          19
          20
             ax[1].set title('Testing Accuracy & Loss')
          21
             ax[1].legend()
             ax[1].set xlabel("Epochs")
         22
             ax[1].set ylabel("Training & Validation Loss")
          23
          24
             plt.show()
```



WARNING:tensorflow:From <ipython-input-29-e74f08e66a06>:1: Sequentia l.predict\_classes (from tensorflow.python.keras.engine.sequential) is deprecated and will be removed after 2021-01-01.

Instructions for updating:

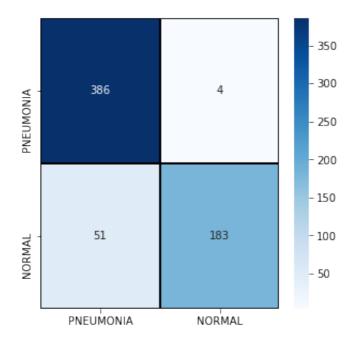
Please use instead:\* `np.argmax(model.predict(x), axis=-1)`, if yo ur model does multi-class classification (e.g. if it uses a `softm ax` last-layer activation).\* `(model.predict(x) > 0.5).astype("int32")`, if your model does binary classification (e.g. if it uses a `sigmoid` last-layer activation).

```
In [30]: fidation_report(y_test, predictions, target_names = ['Pneumonia (Class
```

	precision	recall	f1-score	support
Pneumonia (Class 0)	0.88	0.99	0.93	390
Normal (Class 1)	0.98	0.78	0.87	234
accuracy			0.91	624
macro avg	0.93	0.89	0.90	624
weighted avg	0.92	0.91	0.91	624

```
In [33]: 1
2
olôr = 'black' , linewidth = 1 , annot = True, fmt='',xticklabels = lal
```

#### Out[33]: <AxesSubplot:>



#### **Result:**

Model2 without dropout layer and 6 convolutional layers has an accuracy of 91%, and a recall of 99% on Pneumonia. The no of false positives in this model is less than the first one.

# **Building a CNN3 Model:**

### with RMSprop optimizer:

```
In [49]: ildign the CNN2 model
el3 = Sequential()
ding the first layer
ep1: Convolution
el3.add(Conv2D(32,(3,3),strides =1,padding='same',activation ='relu',in
ep2:Pooling
el3.add(MaxPool2D((2,2)))

8
ding a second convolutional layer
el3.add(Conv2D(64,(3,3),strides =1,padding='same',activation ='relu'))
de1.add(Dropout(0.2))
```

```
oll 2nh
el3.add(MaxPool2D((2,2)))
 14
ding a third convolutional layer
ell3.add(Conv2D(128,(3,3),strides =1,padding='same',activation ='relu')
ell7add(Dropout(0.3))
ol Bng
ell 3.add(MaxPool2D((2,2)))
 20
e@3: Flattening
ella.add(Flatten())
ell3.add(Dense(units =256,activation = 'relu'))
ella.add(Dropout(0.3))
 25
e@4: Full connection
el3.add(Dense(units=1,activation ='sigmoid'))
maBing the CNN
ell.compile(optimizer = "rmsp",loss = binary crossentropy',metrics = ['d
splay model summary.
eB3.summary()
```

Model: "sequential 4"

Layer (type)	Output	Shape	Param #
conv2d_14 (Conv2D)	(None,	150, 150, 32)	320
max_pooling2d_14 (MaxPooling	(None,	75, 75, 32)	0
conv2d_15 (Conv2D)	(None,	75, 75, 64)	18496
max_pooling2d_15 (MaxPooling	(None,	37, 37, 64)	0
conv2d_16 (Conv2D)	(None,	37, 37, 128)	73856
max_pooling2d_16 (MaxPooling	(None,	18, 18, 128)	0
flatten_3 (Flatten)	(None,	41472)	0
dense_10 (Dense)	(None,	256)	10617088
dropout_8 (Dropout)	(None,	256)	0
dense_11 (Dense)	(None,	1)	257
T. 1. 710.017			

```
Total params: 10,710,017
Trainable params: 10,710,017
Non-trainable params: 0
```

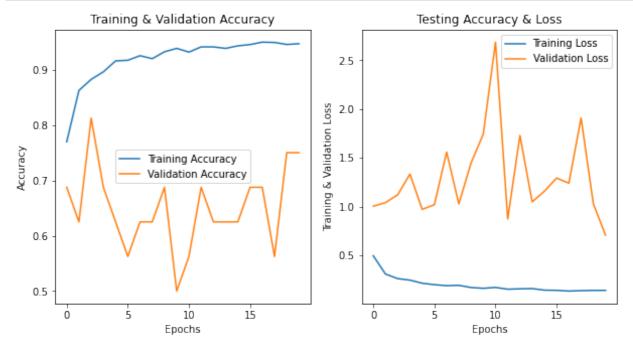
```
4# end timer
5end = datetime.datetime.now()
6elapsed = end - start
7print('Training took a total of {}'.format(elapsed))
```

```
Epoch 1/20
.4969 - accuracy: 0.7697 - val loss: 1.0052 - val accuracy: 0.6875
Epoch 2/20
.3099 - accuracy: 0.8625 - val loss: 1.0414 - val accuracy: 0.6250
Epoch 3/20
29 - accuracy: 0.8825 - val loss: 1.1217 - val accuracy: 0.8125
Epoch 4/20
77 - accuracy: 0.8961 - val loss: 1.3339 - val accuracy: 0.6875
Epoch 5/20
.2159 - accuracy: 0.9158 - val loss: 0.9723 - val accuracy: 0.6250
Epoch 6/20
08 - accuracy: 0.9170 - val loss: 1.0200 - val accuracy: 0.5625
Epoch 7/20
06 - accuracy: 0.9252 - val_loss: 1.5594 - val_accuracy: 0.6250
Epoch 8/20
.1943 - accuracy: 0.9199 - val loss: 1.0280 - val accuracy: 0.6250
Epoch 9/20
.1719 - accuracy: 0.9323 - val loss: 1.4464 - val accuracy: 0.6875
Epoch 10/20
.1637 - accuracy: 0.9385 - val loss: 1.7432 - val accuracy: 0.5000
Epoch 11/20
.1727 - accuracy: 0.9317 - val_loss: 2.6855 - val_accuracy: 0.5625
Epoch 12/20
.1546 - accuracy: 0.9411 - val loss: 0.8739 - val accuracy: 0.6875
Epoch 13/20
.1583 - accuracy: 0.9411 - val loss: 1.7292 - val accuracy: 0.6250
Epoch 14/20
.1609 - accuracy: 0.9385 - val loss: 1.0491 - val accuracy: 0.6250
Epoch 15/20
.1448 - accuracy: 0.9431 - val loss: 1.1583 - val accuracy: 0.6250
Epoch 16/20
.1430 - accuracy: 0.9454 - val loss: 1.2932 - val accuracy: 0.6875
```

### In [51]:

```
#Evaluate on test:
print("Loss of the model is - " , model3.evaluate(x_test,y_test)[0]
print("Accuracy of the model is - " , model3.evaluate(x_test,y_test)
```

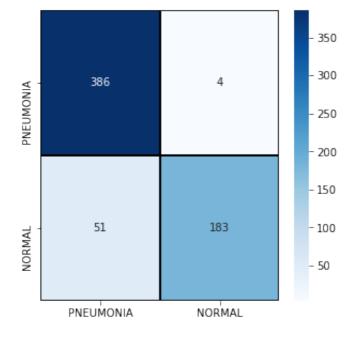
```
In [52]:
             #plotting training& validation accuracy, Traning & validation loss
             import matplotlib.pyplot as plt
           2
           3
             epochs = [i for i in range(20)]
             fig , ax = plt.subplots(1,2)
             train acc = history.history['accuracy']
             train loss = history.history['loss']
           7
             val acc = history.history['val accuracy']
             val loss = history.history['val loss']
          9
             fig.set size inches(10,5)
          10
             ax[0].plot(epochs , train_acc , label = 'Training Accuracy')
          11
             ax[0].plot(epochs , val_acc , label = 'Validation Accuracy')
         12
         13
             ax[0].set title('Training & Validation Accuracy')
          14
             ax[0].legend()
             ax[0].set xlabel("Epochs")
         15
             ax[0].set_ylabel("Accuracy")
          16
         17
             ax[1].plot(epochs , train loss , label = 'Training Loss')
         18
             ax[1].plot(epochs , val loss , label = 'Validation Loss')
          19
          20
             ax[1].set title('Testing Accuracy & Loss')
          21
             ax[1].legend()
             ax[1].set xlabel("Epochs")
         22
             ax[1].set ylabel("Training & Validation Loss")
          23
          24
             plt.show()
```



		precision	recall	fl-score	support
Pneumonia (C	Class 0)	0.88	0.99	0.93	390
Normal (C	Class 1)	0.98	0.78	0.87	234
ā	accuracy			0.91	624
	acro avg	0.93	0.89	0.90	624
weigh	nted avg	0.92	0.91	0.91	624

```
In [55]:  #Plotting confusion matrix.
2  plt.figure(figsize = (5,5))
3  sns.heatmap(cm,cmap= "Blues", linecolor = 'black' , linewidth = 1
```

#### Out[55]: <AxesSubplot:>



#### **Results:**

#### Comparing scores of the models.

### **Conclusion:**

- 1. Activation function used was Relu throughout except for the last layer Sigmoid was used as it is a binary classification problem.
- 2. Dropout layers have been added to reduce overfitting.
- 3. Optimizer for the first 2 models is 'adam', for the last model3 'rmsprop' has been used.
- 4. Loss function used is binary crossentropy.
- 5. Max no of epochs used is 20, with batch size of 32.

### **Future Improvements**

- 1. Training selected models with a a higher no of epochs to try to reach convergence.
- 2. Gathering more data for a better model.
- 3. Testing this data on different pretrained models would lead to significant improvement.
- 4. This work can be extended to detect and classify X-Ray images with lung cancer & Pneumonia.
- 5. If model's recall and accuracy scores are good it can surely help in reducing patient wait times.

```
In [ ]: 1
```