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Covid19 and Pneumonia Detection Using Deep-Learning

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Abstract: It became clear that humanity must learn to live with and adapt to any pandemic such as Covid - 19, especially in light of the fact that the vaccines currently in development do not prevent the infection but only lessen the intensity of the symptoms. It is crucial to diagnose both pneumonia and COVID-19 since they both have an impact on the lungs. In this study, the automatic detection of the Coronavirus disease was carried out using a data-set of X-ray pictures from patients with common bacterial pneumonia, confirmed Covid-19 disease, and normal occurrences. The current approach for detection and diagnosis of COVID-19 is the RT-PCR and rapid test as known rapid test is not so effective and RT-PCR is time-consuming and, in lots of instances, no longer less expensive as a consequence the development of new low-price rapid tests of diagnostic gear to aid medical evaluation is needed.

The study's objective is to assess the effectiveness of cutting-edge convolutional neural network designs for medical picture categorization that have been recently suggested. In particular, the Transfer Learning method was utilised. Transfer learning makes it possible to detect many problems in small collections of medical image data, frequently with outstanding outcomes. The data sets used in this investigation are a collection of 5144 X-ray images, including 460 images with verified Covid-19 illness, 3418 photos with confirmed common bacterial pneumonia, and 1266 images of healthy conditions. The information was gathered from X-ray pictures that were accessible in public medical repositories. According to the data, Deep Learning combined with X-ray imaging may be able to identify important biomarkers for the Covid-19 disease. CNN achieved the highest accuracy 99.49% and specificity, followed by VGG-16 at 67.19% and densenet at 91.94

Index Terms: covid - 19, Pneumonia, CNN, VGG - 16, Densenet, Accuracy

I. INTRODUCTION

Although COVID-19 is an acutely resolving illness, it has the potential to be fatal. Due to extensive alveolar damage and developing respiratory failure, severe illness may cause death when it first manifests. Countries may benefit from the early and automatic diagnosis of Covid-19 for the timely referral of the patient to quarantine and the quick incubation of dangerous cases. In specialised hospitals, and the disease's spread is tracked. Although the diagnosis procedure has become reasonably quick, the financial problems brought on by the expense concerns both states and patients, particularly in nations, regarding diagnostic examinations. The X-rays from healthy instances are now more frequently available to the public than they were in 2020, although additionally from individuals who have Covid-19. As a result, we can examine the medical photographs and recognise any patterns that could result in the condition being automatically diagnosed.

Pneumonia is a condition when the lungs' air sacs fill with fluid, causing inflammation in the lungs as a result. This leads to other problems like chills, weariness, and fever as well as breathing difficulties, coughing, and chest pain. The common cold and the flu are both causes of pneumonia. However, bacterial infections and viral infections can also be problematic at times. This is the cause; one of the factors cited as contributing to pneumonia is COVID 19.

The distinction between COVID 19-infected pneumonia and regular pneumonia has recently been the subject of numerous investigations and experiments conducted in laboratories by medical professionals. In order to illustrate how both have some significant differences, IDSA has developed a thorough knowledge using CT scan images and other sources. Therefore, it is essential to diagnose Pneumonia and COVID.

Deep learning applications seem to have emerged throughout the last five years at the perfect time. Automatically identifying features in photographs and classifying them is a major focus of the "Deep Learning" class of machine learning techniques. Its main applications are in tasks involving the classification of medical images and item identification. When it comes to using artificial intelligence to mine, analyse, and uncover patterns in data, deep learning and machine learning are well-established fields of research. The contributions made by those disciplines to clinical decision making and computer-aided systems are getting harder to recapture as new data become accessible.

Deep neural networks, also known as deep learning, are artificial neural networks (ANN) with several layers. Due to its capacity to manage massive volumes of data, it has emerged during the last few decades as one of the most effective tools and has seen significant literary success. In a number of applications, including pattern recognition, deeper hidden layers are already beginning to perform better than conventional methods. One of the most popular deep neural networks is the convolutional neural network (CNN). Convolutional Neural Networks have made groundbreaking discoveries over the past 10 years in a variety of pattern recognition-related fields, including speech recognition and picture processing. The reduction of ANN's parameter count is CNNs' most advantageous feature. The most crucial premise regarding issues that CNN solves is that they shouldn't include features that are spatially dependent. This is because larger models may be used to handle sophisticated tasks that were not conceivable with classic ANNs. In other words, we don't need to focus on where the faces are in the photographs while using a face detection tool, for instance. The only thing that matters is finding them, regardless of where they are in the provided photographs. Obtaining abstract characteristics when input propagates toward the deeper layers is another crucial component of CNN.

II. RELATED WORK

The author investigated a number of well-known pretrained deep CNN models in a transfer learning setup. For the purpose of identifying COVID-19 from chest X-ray images. Different setups were tested using the two separate publically accessible datasets either alone or jointly. Different setups were tested using the two different publically accessible datasets either alone or together.

Limitation: The COVID-19 vs. non-COVID-19 chest X-ray classification by VGG models was unsuccessful. The COVID-19 vs. non-COVID-19 chest X-ray classification by VGG models was unsuccessful. (from reference [6])

To assess and contrast the created models, three distinct experiments are conducted in accordance with three preprocessing approaches. The objective is to assess the effects of data preprocessing on the outcomes and how well they can be explained. Similar to this, a comprehensive investigation of various variability problems that could potentially ruin the system and its impacts is carried out. The methodology used yields a classification accuracy of 91.5 percent, with an average recall of 87.4 percent for the weakest but most explicable experiment.

Limitation: In this approach, manual methods based on visual observation of the images were used in earlier CT studies of COVID-19 to test the infection extent evaluation. (from reference [1]).

A proposed approach for analysing chest X-ray pics has been created to come across COVID-19 for binary instructions with an accuracy of 78 percentage and validation accuracy of 80 percent, wherein the loss is more or less zero.15 percent. This method applies convolution 2D techniques to COVID-19 open supply datasets which can be available at GitHub and Kaggle

Limitation: massive dataset working with GPU taking into account many more attributes to evaluate for high computational speed, performance, and effective deep learning approaches implementation. (from reference [7])

A streaming diagnosis based on a deep learning-based retrospective analysis of laboratory data in the form of chest X-rays is required in this COVID-19 pandemic situation. In this paper, a method to detect COVID-19 using deep learning to assemble medical images was proposed.

Limitation: For the datasets used to determine if a person is contagious or not, classification techniques must be applied. (from reference [2])

The suggested approaches were validated using a dataset that was specifically retrieved for this study. Despite the poor quality of the chest X-ray images that are a drawback of portable equipment, the proposed approaches produced global accuracy values of 79.62 percent, 80.27 percent, and 79.86 percent, respectively. This allowed a reliable analysis of portable radiographs to support clinical decision-making. Limitation: Although the decision-making accuracy is lower, the results show that the proposed technique and the tested approaches permit a robust and reliable analysis. (from reference [3])

In this article, CNNs are briefly introduced, along with recently released papers and innovative methods for creating these amazingly outstanding image recognition models. The introduction makes the assumption that you are already familiar with the basics of ANNs and machine learning.

Limitation: In contrast to other artificial neural network types, convolutional neural networks concentrate on a specific type of input rather than the entire problem domain. (from reference [5])

Textual clinical reports were divided into four categories in this study using conventional and ensemble machine learning techniques. These features were applied to conventional and group machine learning classifiers.

Limitation: By adding more data, models' effectiveness can be increased. In order to determine whether men or women are more likely to contract the disease, it can be categorised based on gender. Deep learning approaches can be applied in the future, but more feature engineering is required for improved outcomes. [4]

In this study, the author present a unique Support Vector Regression approach to analyse five distinct tasks associated with novel coronaviruses. To improve classification accuracy in this work, supported vectors are also used in place of a simple regression line. Limitation: The encouraging outcomes show its inferior superiority in terms of both efficiency and accuracy..(from reference[8])

III. PROPOSED METHOD

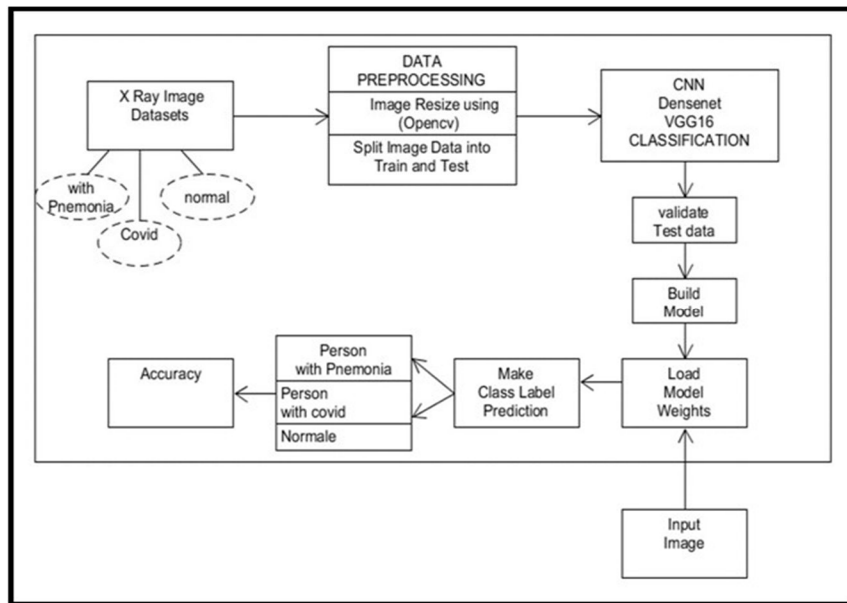


Fig. 1: Proposed Block Diagram

A. Data Collection

The process of gathering and coordinating information from an unlimited number of various sources is known as data collection. In order to process the data after the statistics have been collected, the information is recorded as an image (chest x-ray). A dataset in machine learning is, to put it simply, a collection of data samples that can be analysed and forecasted by a computer as a single entity. This implies that the data gathered should be uniform and understandable because machines don't perceive data in the same manner that people do. A good dataset should also adhere to strict quality and quantity standards. The dataset should also be relevant and evenly distributed for a rapid and simple training procedure. For our work, we created three classes of chest X-ray image datasets (normal people, pneumonia patient, and COVID - 19 Patients).

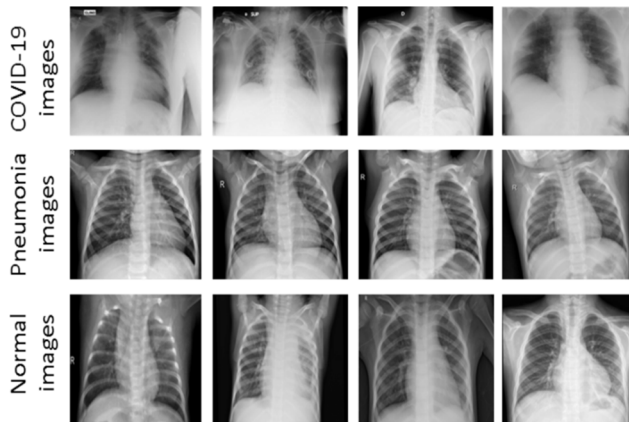


Fig. 2: Data set with three classes

It has 5144 total images, of which 460 are covid-19, 1266 are images of people who are healthy or normal, and 3418 are images of people who have pneumonia. Covid-19 is a recent infection, thus there aren't many pictures of it. The data set is obtained from the Kaggle website and divided into two portions: 20% for validation and 80% for training.

B. Data Preprocessing

Preprocessing is a data mining technique used to transform raw data into a format that is appropriate and useful. The actions needed for data preprocessing are:

1) *Data Cleaning*: The data may contain many irrelevant and missing pieces. Data cleaning is done in order to control this portion. It involves handling noisy, missing, and other types of data.

2) *Data Transformation*: This procedure is used to turn the data into forms that are suitable for the data mining process. It involves the following actions:

a) *Normalization*: To get the values of the data in the specified range, normalisation is done.

b) *Attribute Selection*: The set of already provided attributes is used to construct entirely new attributes.

c) *Discretization*: Restoring the raw values of numerical properties at interval levels is done.

d) *Concept Hierarchy Generation*: Based on hierarchy, the qualities are changed from a low level to a high level.

3) *Data Reduction*: Since managing large amounts of data requires a certain approach, data mining. It was challenging to assess in these situations while working with a significant amount of data. So we used a data reduction strategy to get away from this. Reduced expenses and increased storage efficiency were the goals of the reduction.

C. Split Data

In order to evaluate how well the machine learning algorithm is working, data is split into training and testing sets.

1) *Train Data-set*: treated to bring relevance to the model.

2) *Test Data-set*: treated to assess the relevant model.

The major goal is to assess the model's performance using the most recent data, i.e., data that were not utilised in the model's training. When there are several data sets gathered, the train-test methodology is appropriate.

The size of the training and testing data sets is a crucial configuration constant for the course of action. For either the training or testing data sets, it is often shown as a percentage between 0 and 1. For instance, if the size of the training set is 0.60 (60 percent), the testing set will receive the remaining 0.40 (40 percent). For dividing data sets, there is no ideal split %. Taking into account the needs of our project, we choose a percentage to divide the data sets by.

3) Cost of computation for training a model

4) Computed costs for model evaluation

5) Representation of the training set's behaviour

6) Behaviour of the testing set's representation

We took into account the aforementioned elements when choosing the percentage to divide the data into training and testing sets for this project. We used the Python machine learning toolkit scikit-learn, which provides a training and testing split assessment application. The "test size" option, which accepts an input of the number of rows or a percentage of the size of the data set between 0 and 1, can be used to describe the split size.

D. Algorithms

The deep learning algorithm is a method for the system of AI capabilities to carry out the activities, typically by anticipating the values as output from previously provided data as input.

The important actions of algorithms of deep learning is image classification.

E. Transfer learning with CNNs

Transfer learning is a method that trains a CNN from start to perform a different but related job using new data, often from a smaller population. Transfer learning makes advantage of the data-mining knowledge produced by a CNN.

In this deep learning process, a CNN is first trained using big datasets for a particular objective (like classification). The availability of data for the first training is the most crucial component for training to be successful since CNN may discover critical properties (features) of the image. The suitability of this model for transfer learning depends on the CNN's capacity to recognise and extract the most exceptional visual characteristics. The processing of a brand-new collection of images of a different kind and the use of the CNN to extract features based on its experience with feature extraction since its first training constitute the following phase.

F. CNN

Convolutional neural network (CNN) techniques are typically applied to designs with a large number of training layers. Weights and bias are the two factors for every layer. Later layers search for mid-level and high-level features such as structures, objects, and shapes, while earlier layers concentrate on low-level features like corners, edges, and lines. Using the prediction result, it is then possible to calculate the loss cost to the ground truth labels. In order to compute each parameter's gradient based on the loss cost, the backward stage secondly uses chain rules. For the future gradient-based forward computation, every parameter has been updated and prepared. Following several cycles in both the forward and backward stages, network learning might come to an end.

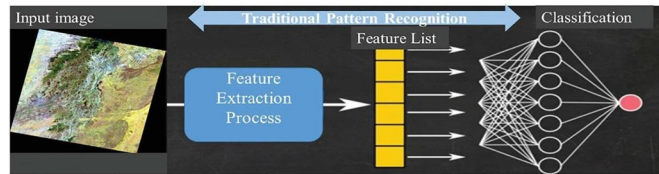


Fig. 3: Layers of CNN

It would be advantageous to define the term since it is frequently used to refer to mathematical notions and ideas connected to the feature transformation strategy and procedure. One of the starting functions in mathematics, most notably algebraic topology, is turned into a third function by the mathematical procedure known as convolution. This third function is frequently viewed as a transformed version of the two initial functions in mathematics, u and v . The convolution is a distinct function that is frequently represented by the letters u and v and discussed by (Hirschman and Widder, 2017) in the context of two real or complex functions, u and v .

$$(uvx)(t) = \int v(t-d)u(d)dt \tag{1}$$

G. ReLU Layer

The layer is known as Rectified Linear Units. This layer of neurons employs the loss function or non-saturating non-linearity function: $f(x) = \max(0, x)$

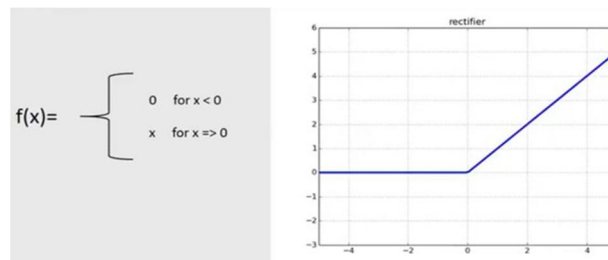


Fig. 4: Relu Activation Function Graph

Rectified Linear Unit (ReLU) transform functions only turn on a node if the input value exceeds a specific threshold. When the information rises above a threshold, the output changes from zero while the data is below zero. It and the dependent variable are related linearly.

1) *Pooling Layer*: Condensing or reducing the spatial dimensions of feature map-derived data is its mission. The most common type of pooling is maximum pooling because of its enhanced convergence and speed. Average pooling and L2-norm pooling are the other two types (Scherer et al., 2010). In essence, this requires a filter, commonly of size 2×2 , and an equal-length stride. With a filter that convolves after being applied to the input volume, it produces the highest number possible in each subregion.

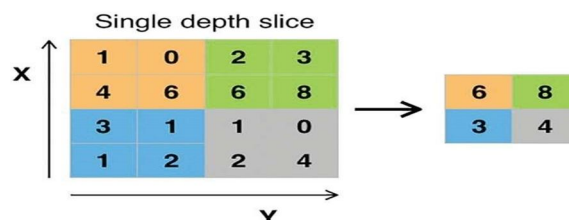


Fig. 5: Max Pooling Layer

H. Fully connection layer

An input vector of numbers is provided to this layer. The phrase "fully connected" describes a system in which every input is coupled with every output. In the CNN procedure, it often comes after the final pooling layer. 90% of the CNN's parameters are located in fully linked layers, which function similarly to a traditional neural network. This layer basically receives the output of the previous pooling layer and creates an N-dimensional vector, where N is the total number of classes the computer can choose from. We can use it to feed a vector with a specific length forward into the neural network. Additionally, the result is a vector of numbers. This

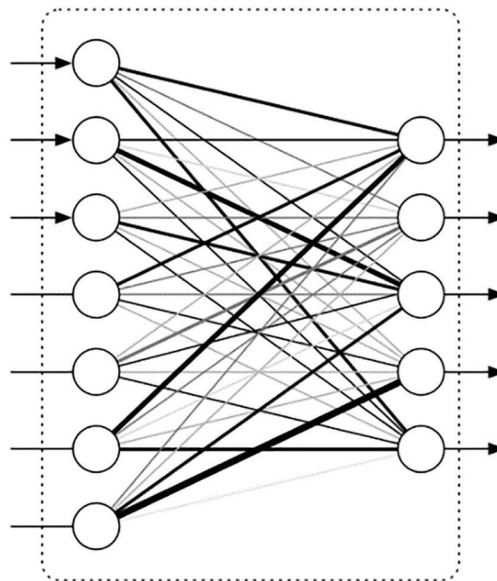


Fig. 6: Fully connected layer

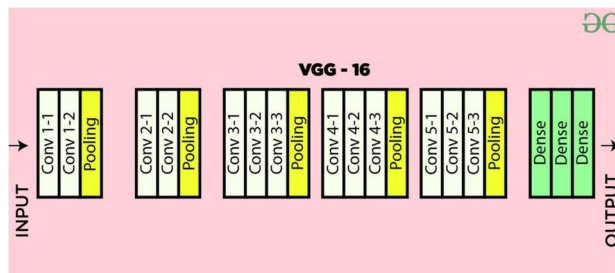
simulates higher-level reasoning in which all potential routes from the input to the output are taken into account. After passing through two layers of convolution, relu, and pooling, and being converted into a single file or a vector, take the reduced image and place it in the single list.

- 1) *Loss Layer*: Loss layer computes a value that represents the model's performance by using functions that take the model's output and the target into account. It may serve two primary purposes.:
 - a) Forward (input, target): based on input and target value, estimates loss value.
 - b) Backward (input, target): identifies the criterion, calculates the gradient of the loss function associated with it, and returns the outcome.
 - c) In a CNN, the back propagation principle is employed to determine the gradient of the loss function, which determines the cost related to a particular state.
- 2) *VGG - 16*:

I. VGG - 16 Architecture

The term "ConvNets" is frequently used to refer to convolutional neural networks, a subset of artificial neural networks. Input, output, and multiple hidden layers are the components of a convolutional neural network. One of the most effective computer vision models available right now is the CNN(Convolutional Neural Network) variant known as VGG16. In order to analyse the networks and deepen the model, the developers developed an architecture with incredibly small (3x 3) convolution filters, which was a substantial advance over the state-of-the-art setups. Around 138 trainable parameters were created with the depth increased to 16-19 weight layers. An input image with dimensions is sent to the network (224, 224, 3). In the first and second layers, there is the same padding and 64 channels with a 3*3 filter size. Following a stride (2, 2) max pool layer, two layers have convolution layers with a 128 filter size and a filter size (3, 3). Max-pooling stride(2, 2) layer with the exact same properties as the layer before follows. Then, 256 filters are spread out over 2 convolution layers with 3 and 3 filter widths.

A max pool layer comes next, then two sets of three convolution layers. With the same spacing and 512 filters per filter (3, 3). Then, this image is subjected to the stack of two convolution layers. We employ 3*3 filters in these convolution and max-pooling layers as opposed to 7*7, ZF-11*11, and AlexNet filters. Additionally, several of the layers change the number of input channels by using 1*1 pixels. A 1-pixel padding is offered after each convolution layer to prevent the spatial characteristic of the image



.Fig. 7: VGG - 16 Architecture

As Convolution and max-pooling layers were added to the stack, and the result was a (7, 7, 512) feature map. A feature vector with the value 1, 25088 is produced by flattening this result. There are then three fully connected layers: the first layer uses the most recent feature vector as input and outputs a vector with a size of (1, 4096); the second layer also does so; the third layer, however, also produces a vector with a size of (1, 1000), which is used to implement the softmax function to divide the data into 1000 categories. Every hidden layer has a ReLU activation function. ReLU is more efficient since it encourages faster learning and reduces the possibility of error.

J. Densenet

One of the most recent advancements in neural networks for visual object detection is called DenseNet. Despite their apparent closeness, ResNet and DenseNet have some important differences. While DenseNet uses an additive approach to combine the previous and subsequent levels, ResNet concatenates the layers.

Composition Layer Pre-Activation For each composition layer that produces feature maps with k channels, Batch Norm (BN), ReLU, and 33 Conv are executed. For instance, X0, X1, X2, and X3 may all be altered to X4. The concept for this was generated by Pre-Activation ResNet.

K. DenseNet-B (Bottleneck Layers)

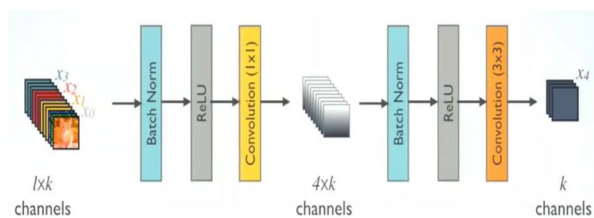


Fig. 8: Graph of accuracy and training losses

To reduce the model complexity and size, BN-ReLU-1x1 Conv is done before BN-ReLU-3x3 Conv.

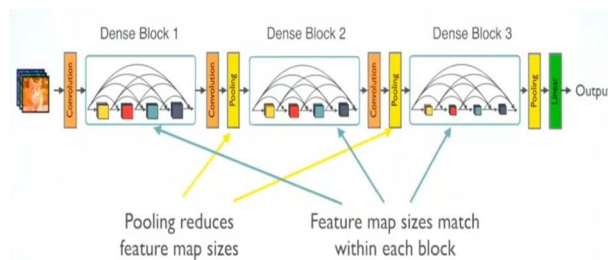


Fig. 9: Multiple Dense Blocks with Transition Layers

1) *Multiple Dense Blocks with Transition Layers:* Multiple Dense Blocks In addition to 2 x 2 common pooling, 1 x 1 Conv is used as the transition layers between neighbouring dense blocks. The dense block has a set of feature maps, each of which is the same size, making it easy to concatenate them. It is easy to concatenate the dense block's feature maps together due to their similar sizes. Upon completion of global average pooling at the end of the last dense block, a softmax classifier is linked.

L. DenseNet-BC (Further Compression)

The transition layer produces m output feature maps, with α being the compression factor, if there are m feature-maps in a dense block. On each transition level when $\alpha=1$, a fixed number of feature-maps are present. DenseNet-C, also known as DenseNet, had a value of 1 and was equal to 0.5 throughout the trial. When the bottleneck layer and the transition layer with 1 are both used, the model is referred to as DenseNet-BC. Currently, DenseNets with/without B/C, various L layers, and various k growth rates are also being trained.

M. Prediction

The result of the algorithm, which shows if a specific illness as covid-19 or pneumonia can be predicted based on the data sets, has been trained on a genuine data-set and reinforced to recent data.

N. Accuracy

The study used to show which machine learning algorithm's model is best at identifying connections and patterns between different variables in data sets is known as the model accuracy.

O. Confusion Matrix

It is a 2x2 binary classification matrix with real values on one axis and predicted values on the other. we introduce two concepts: false positives and false negatives.

		ACTUAL	
		Negative	Positive
PREDICTION	Negative	TRUE NEGATIVE	FALSE NEGATIVE
	Positive	FALSE POSITIVE	TRUE POSITIVE

Confusion Matrix

Fig. 10: confusion matrix

- 1) True Positive (TP) — model successfully predicts the positive class (prediction and actual both are positive).
- 2) True Negative (TN) — model predicts the negative class accurately (prediction and actual both are negative).
- 3) False Positive (FP) — incorrect prediction of the negative class is made by the model. (predicted-positive, actual-negative).
- 4) False Negative (FN) — Model predicts the positive class inaccurately. (predicted-negative, actual-positive).

$$TPR = \frac{TP}{Actual\ Positive} = \frac{TP}{TP + FN}$$

$$FNR = \frac{FN}{Actual\ Positive} = \frac{FN}{TP + FN}$$

$$TNR = \frac{TN}{Actual\ Negative} = \frac{TN}{TN + FP}$$

$$FPR = \frac{FP}{Actual\ Negative} = \frac{FP}{TN + FP}$$

- a) *Precision*: The precision ratio is the sum of the true positives and false positives. i.e. What proportion of all the positive predictions that are actually positive.

$$Precision = \frac{TP}{TP + FP}$$

The precision number ranges from 0 to 1.

- b) *Recall*: Recall examines the number of erroneous negatives that were included in the prediction process rather than the number of false positives the model predicted, i.e., what proportion of all positives are anticipated to be positive. It is equivalent to TPR (true positive rate).

$$Recall = \frac{TP}{TP + FN}$$

- c) *F1 - Score*: It is the precision and recall harmonic mean. It accounts for both false positives and false negatives. Consequently, it works well with an unbalanced dataset.

$$F1\ score = \frac{2}{\frac{1}{Precision} + \frac{1}{Recall}} = \frac{2 * (Precision * Recall)}{(Precision + Recall)}$$

Recall and precision are given the same weight in the F1 score. We can assign varying weights to recall and precision in a weighted F1 score. Recall and precision are given varying weights in different tasks, as was covered in the preceding section.

$$F_{\beta} = (1 + \beta^2) * \frac{(Precision * Recall)}{(\beta^2 * Precision) + Recall}$$

Beta measures the frequency with which recall outweighs precision. The value of Beta is 2 if recall matters twice as much as precision.

IV. RESULTS AND ANALYSIS

The results, which were acquired using a variety of algorithms including CNN, VGG-16, and densenet, are compared, examined, and predicted; they are then described in more detail below.

A. Results Discussion

We performed the classification of covid - 19, pneumonia infection utilizing python 3 on jupyter notebook, sklearn models and predictions and utilized the CNN algorithms namely CNN, VGG - 16, Dense net, and got the accuracy and loss of the models and performed comparison between these algorithms and obtained the below results.

B. Result of CNN Algorithm

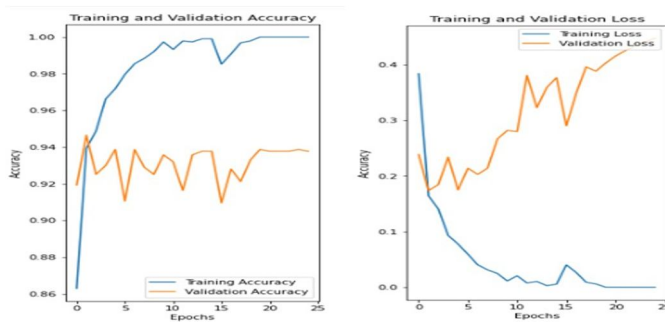


Fig. 11: Graph of Accuracy and Losses for CNN

The trending graph that shows how training and validation accuracy change with epoch count is shown in the figure above. The graph suggests that there is a significant increase in training accuracy as soon as the number of epochs rises, based on detailed examination of the data. 32 batches of 25 epochs each are used to train the model. As a result, training across 25 of these epochs produced results that were optimal, with high accuracy of 100% and a minimal loss of 4.0362e-05 for training and a similar accuracy of 93.7 % and a loss of 0.44 for validation.

C. Result of VGG - 16 Algorithm

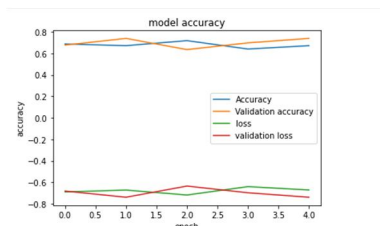


Fig. 12: Graph of Accuracy and Losses for VGG – 16

The VGG -16 model was only trained for 5 epochs because it takes much longer time that the training losses and validation losses both reached negative values, making it clear that the model was overtrained and the over-fitting was evident. Consequently, training for five of these epochs produced results with training accuracy of 67% and validation accuracy of 73%.

D. Result of Dense-net Algorithm

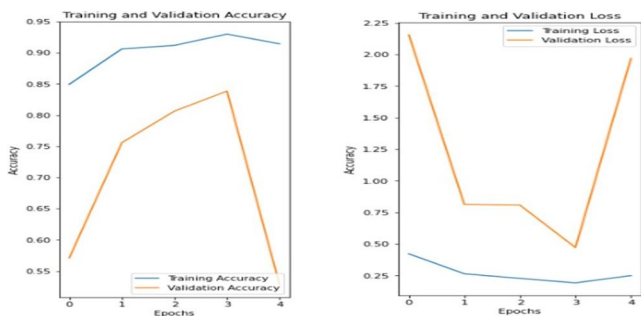


Fig. 13: Graph of Accuracy and Losses for Densenet

Due to the lengthy training process, the Densenet model was only trained for five epochs. The training phase of the model is conducted across 5 epochs, with a batch size of 16 for each epoch, for a total of 100 steps. Consequently, training across 25 of these epochs produced optimum results, with good accuracy of 94 percent and a low loss of 0.22 for training and a similar accuracy of 52 percent and a loss of 1.97 for validation. From the obtained confusion matrix we can determine performance of the classification models for a given set of test data by calculating the precision, recall and F1 - Score values.

E. Confusion matrix

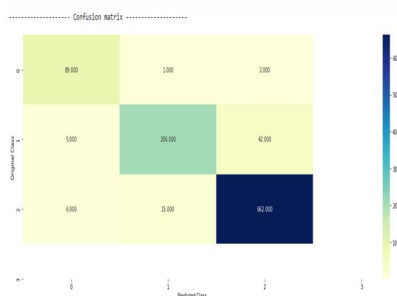


Fig. 14: Confusion Matrix

TABLE I: Classification Report for Densenet

	Covid	Normal	Pneumonia
Precision	0.89	0.92	0.93
Recall	0.96	0.81	0.96
F1 Score	0.92	0.86	0.94

F. Algorithm comparison

The comparison of all three algorithms is provided in the table above so that we can determine which model is ideal for creating web applications. and can also assess how each model behaves.

TABLE II: Training Accuracy

No of epoch	CNN	VGG-16	Densenet
1	86.30	68.75	84.94
2	93.93	67.19	90.75
3	94.87	71.88	91.44
4	96.62	64	91.75
5	97.96	67	91.06
6 - 10	99.73	overfitting and longtime	longtime
11 - 25	100	overfitting and longtime	longtime

TABLE III: Validation Accuracy

No of epoch	CNN	VGG-16	Densenet
1	91.93	67.71	57.1
2	94.65	73.96	75.58
3	92.51	63.54	80.64
4	93	69.79	83.85
5	93.87	73.96	52.82
6 - 10	93.58	overfitting and longtime	longtime
11 - 25	93.77	overfitting and longtime	longtime

TABLE IV: Training Losses

No of epoch	CNN	VGG-16	Densenet
1	38.36	- 69	42.18
2	16.43	- 67.28	26.35
3	14.09	-71.87	22.07
4	9.3	-64.06	19.17
5	7.7	-67.19	24.92
6 - 10	1.14	overfitting and longtime	longtime
11 - 25	4.0362e-05	overfitting and longtime	longtime

TABLE V: Validation Losses

No of epoch	CNN	VGG-16	Densenet
1	23.86	- 68.21	71.79
2	17.38	- 73.96	78.05
3	18.44	-63.54	28.04
4	23.39	-69.79	42.15
5	17.48	-73.96	22.59
6 - 10	28.18	overfitting and longtime	longtime
11 - 25	44.76	overfitting and longtime	longtime

From above training and validation accuracy table we observe how the accuracy varies for every epoch. we observe that for the first epoch in training accuracy CNN has around 86% and for further the accuracy increases but in VGG - 16 its around 60% and for further epochs its around 60% as result of over fitting. Dense accuracy also keeps on increasing for epochs but densenet and VGG - 16 whereonly executed for five epochs has its taking loner time.

From above training and validation loss table we can analyze how the losses have reduced for increasing epochs here the CNN exits the very minimal losses(4.0362e-05)hencefrom this comparision its evident that CNN is a best algorithmwhen compared to VGG - 16 and Densenet in our study.

G. Results of web - application

It is clear from the comparison table that the CNN model has good accuracy, which is why a web application was developed for it. The results exhibits how the the developed web application is identifying the X- rays belonging to three different classes covid - 19 , Pneumonia and normal.

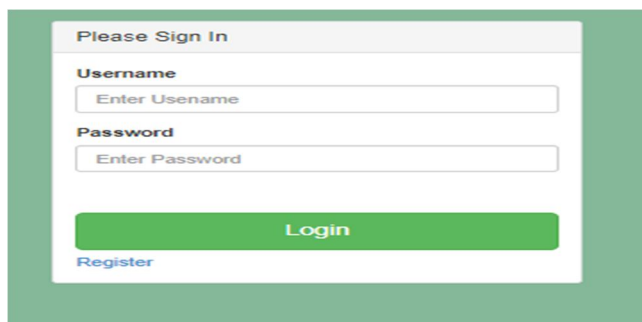


Fig. 15: Front End Developed

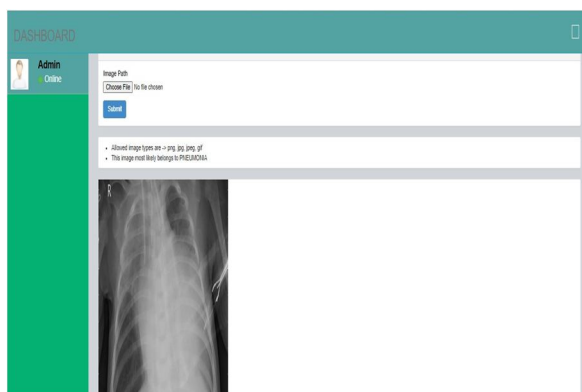


Fig. 16: Dash Board Design

1) *Detection of Covid -19 , Pnuemonia and Healthy chest x- ray:* Here the X - ray images are fed by clicking on choose file in dash board after that we need to select file in which the images are present and upload it once after uploading need to click on submit button once after the submitting image we can observe result whether the given X - ray belong to covidor pneumonia or normal.The same procedure is followed for many iteration to check whether develop web - application is accurate by feeding different types of X - ray images.

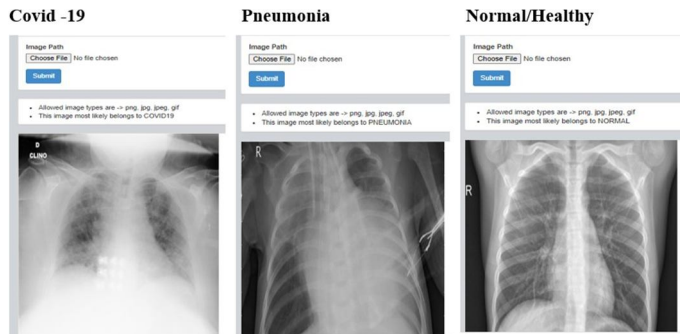


Fig. 17: Iteration 1

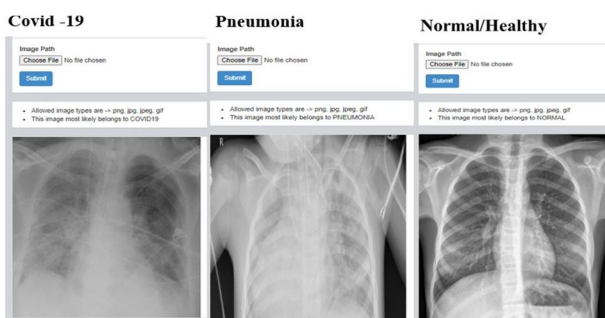


Fig. 18: Iteration 2

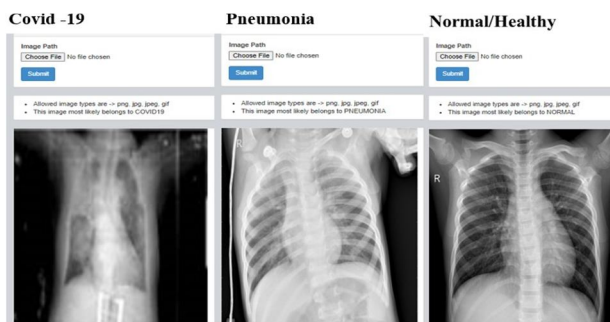


Fig. 19: Iteration 3

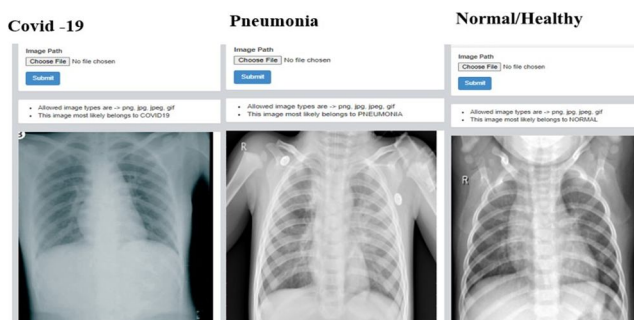


Fig. 20: Iteration 4

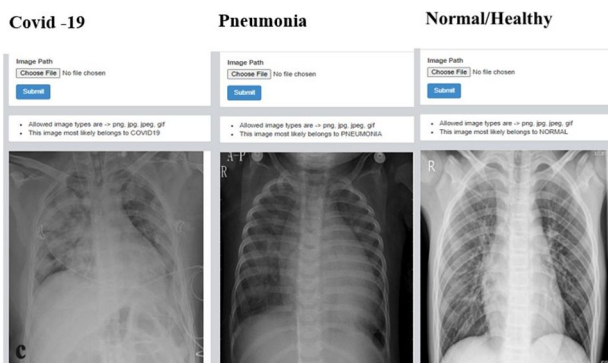


Fig. 21: Iteration 5

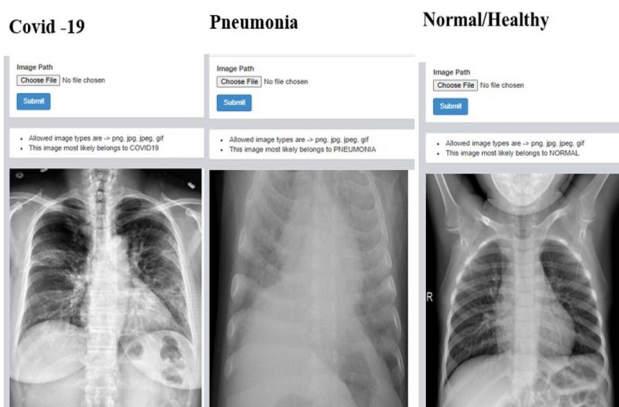


Fig. 22: Iteration 6

V. CONCLUSION

In this study, CNN, dense net, and vgg16 were enhanced to better detect COVID, viral pneumonia, and to differentiate COVID-19 cases from non-COVID-19 cases on chest X-ray pictures. The proposed model eliminates the need for human feature extraction due to its automated nature and end-to-end structure. Less cases were used to train the deep models in the vast majority of earlier research. A variety of example edged images are produced by the proposed model's multi-image augmentation technique, which is supported by first and second order derivative edge operators. The classification accuracy of xray scans and photographs is evaluated using

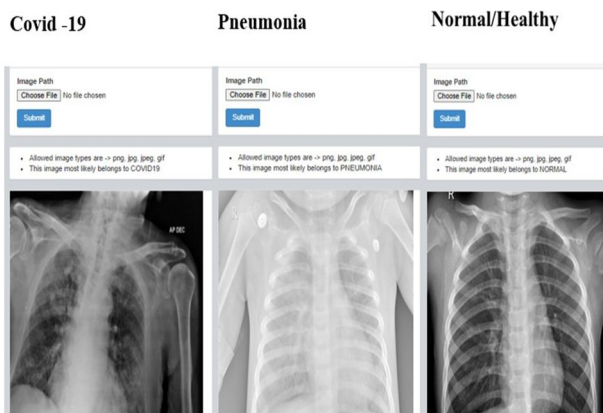


Fig. 23: Iteration 7

CNN that has been trained using these improved images. CNN has a classification accuracy of 100% for X-Ray pictures and photos when it is trained with these improved images. The experimental outcomes were deemed to be quite convincing, and they were helpful for COVID-19 screening on chest X-ray images of individuals who potentially have a corona.

Future research may focus on the early diagnosis of several illnesses, such as pneumonia, bronchitis, and tuberculosis, as well as the COVID-19 of those who are suspected of having a respiratory disorder. Our suggested multi-model ensemble detection strategy has improved compared to the original detection effect for CNN, even though the detection effect for VGG-16 is still insufficient because of over fitting. The lengthy training schedule is another problem. To improve the model's accuracy, each of these elements should be considered. Future study will focus on developing more precise classification systems for the diagnosis of two types of pneumonia caused by bacteria and viruses. The CNN-based version is a promising way to use X-rays to identify the disease, according to the description that was previously provided.

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