

Classification of Covid-19 Chest X-Rays Utilizing Deep Convolutional Recurrent Neural Network with RBF Kernel

Pimple Shweta Parshuram¹ , Dinesh Sahu² ¹MTech Student, ²Head and Professor Department of CSE, SRK University, Bhopal, India

Abstract- Governments are able to disrupt the spread of the deadly severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) with the help of early detection and clinical skill. Chest X-ray (CXR) imaging is more accurate for illness categorization and evaluation than reverse transcription-polymerase chain reaction (RT-PCR), yet RT-PCR provides rapid findings. Research into creating a COVID-19 detection toolbox has been sparked by the fast spread of the 2019 coronavirus illness (COVID-19). In order to learn features, deep learning-based approaches like convolutional neural networks (CNNs) need a lot of training data, however recent research has shown that this is the best method for COVID-19 classification. It has been difficult during the epidemic to quickly collect enough training data. The purpose of this research was to propose a novel model for classifying CXR pictures into normal, pneumonia, and COVID-19 using CNNs and deep convolutional recurrent neural networks (DCRNNs). The proposed model outperforms the current pretrained approaches (AlexNet and GoogLeNet) thanks to its eight convolutional layers, four maxpooling layers, and two fully connected layers. DCRNN creates manufactured or counterfeit pictures to make up for a lopsided dataset and furthermore separates profound elements from each picture in the dataset. Also, it widens the informational collection and catches variety's characterizing highlights to further develop speculation. The proposed CNN and the current pretrained methods were prepared and tried on four separate publicly accessible datasets of chest X-beam pictures (Coronavirus X-beam, Coronavirus Chest X-beam, Coronavirus Radiography). After then, the proposed CNN approach was prepared utilizing the previously mentioned four datasets in view of the DCRNN engineered pictures, yielding superior exactness (95.3%, 96.2%, and 99.2%) over the pre-prepared models currently being used. The discoveries overall show that the recommended DCGAN-CNN method is a reasonable choice for exact Coronavirus conclusion.

Key Terms: DCRNN-CNN, AlexNet, GoogLeNet, CXR images.

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I. INTRODUCTION

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Most essential consideration specialists address constant sicknesses the greater part of their time. The drawn-out clinical issues of seven out of each 10 fatalities in the United States are responsible for constant diseases like cardiovascular issues, weight and malignancy. During the earlier decade there were many hereditary variations that may clarify contrasts in rate of persistent sicknesses and pharmacological

response in the individual, giving essential consideration specialists an unrivaled shot at customizing disease expectation and treatment. Also, the cost of such tests has declined rapidly and turn out to be progressively common thus. An assortment of advances ought to be carried out into wellbeing frameworks and clinical cycles to make customized medication completely acknowledged in the Electronic Health Record (EHR) period. They incorporate the wellbeing hazard evaluation (HRA), drug compromise, and physicist choice help. HRAs examination a solid individual by foreseeing sickness affectability and, in specific circumstances, present a suggested hazard the board activity plan. A vital part of HRA is the historical backdrop of the family, which is the most grounded singular indicator for normal persistent issues including diabetes, stroke and cardiovascular diseases and other wellbeing problems, for example, bosom malignancy and colonic disease.

II. BACKGROUND

The current COVID outbreak around the world has been very bad. This is still a big problem around the world, even though medicine and technology have come a long way. Even though medicine has come a long way, the ways we use now are not good enough for finding COVID signs early. People are having trouble making a living now that this illness has spread around the world. CT scan data can now be used by radio testing to find COVID cases. Because of this, the authors of the study talk about how a belief function and convolution neural networks can be used to classify things that could be used to spot the signs of this virus. This method chooses how to group data by first pulling out the important features and then connecting those features with belief maps. This study would put things into groups more accurately than other studies have. With an accuracy of 0.87, an F1 of 0.88, and an area under the bend (AUC) of 0.95, the suggested method is better than the usual deep learning method [1].

By the middle of 2020, the 2019 Covid Sickness (Coronavirus) will have spread all over the world. Robots that can look at computed tomography (CT) pictures and find lung diseases could make the current way of treating Coronavirus a lot more effective. On the other hand, separating infected areas from CT slices can be hard because there is a wide range of contamination quality and there isn't much power difference between healthy and sick tissues. There are also problems with getting a lot of data in a short amount of time, which makes it harder to make a detailed model. To get around these problems, we recommend using a high-tech Coronavirus Lung Contamination Division Deep Network (Inf-Net) to reliably tell the difference between infected areas and chest CT scans. The major level features are put together into a world reference by our Inf-Net using an equal fractional encoder. After that, the limits are shown, and the pictures are made better by using the explicit edge consideration and the implicit switch consideration. Plus, we have a partly controlled split system based on a randomly chosen spread method that works with mostly unstructured data and only a few tagged pictures. Our partially directed framework makes it possible for even more learning to happen and for talks to be of higher quality. A lot of tests using our Coronavirus SemiSeg and real CT volumes show that the suggested Inf-Net works better than most current segmentation models and is better than current performance [2].

A new type of the Covid virus caused the 2019 Coronavirus pandemic. People all over the world are quickly getting the coronavirus. The most accurate way to diagnose Coronavirus is with the opposite transcriptase polymerase chain reaction (RT-PCR) test. But because there aren't many places to do RT-PCR tests, it's hard to get a quick diagnosis of the illness. X-rays and other easily accessible imaging techniques can be used to see the obvious bad effects of the coronavirus. Convolutional neural networks that have already been set up are often used to help PCs find diseases in smaller datasets. This study looks into the possibilities of multi-CNN, which is a group of many already-made CNNs, as a way to automatically find Coronavirus in X-ray pictures. A multi-CNN, a relationship-based highlight selection (CFS) approach, and a Bayes net predictor are used together in this method to predict the presence of Coronavirus. The method was tested on two datasets that were open to the public. The results were good in both cases. With an AUC of 0.963 and an accuracy of 91.16% on the first dataset, which had 453 images of Coronavirus and 497 images of other viruses, the method worked well. The second set had 71 pictures of Coronavirus

and 7 pictures of other viruses. On this set, the method had an AUC of 0.911 and an accuracy of 97.44% [3].

A three-stage Vulnerable Contaminated Recovered Dead (3P-SIRD) model is suggested in this paper to find the best lockdown period for certain geological locations. This will help break the chain of transmission, help the country's economy recover, and help the system fight Coronavirus. The new model is different because it includes hurdles besides just the sickness rate, the thought rate, and the death rate. These include quiet carriers, the kindness of newly infected people, and unlisted Covid infected people who have been passed on. Along with the more basic limits, they make a huge difference in how the model makes sense. During the course of the outbreak, the model looks at how often people are tested and how that frequency changes over time. The suggested 3P-SIRD model is split into three separate stages that take into account how long the illness lasts and how aware the person is of it. As the rate of illness and healing changes from place to place, time is split into several chunks. Testing the model with data from China shows that it works well enough to make predictions that are very close to the real numbers of people sick, people who have recovered from illness, people who have died, and people who are still having symptoms. The model says that the best length of lockdown for China would be 73 days, which is pretty close to the real length of lockdown (77 days). Besides that, the model is used to figure out when the best lockdown time is in India and Italy [4].

That is why this paper look into the long-term effects of the Coronavirus in India after it first showed up in Wuhan, China, in 2019. In order to stop the feast of Coronavirus, India shut down the whole country on March 25, 2020. The Powerless Uncovered Irresistible Recuperated (SEIR) model is used to predict dynamic Coronavirus cases in India. It takes into account the effect of cross-country shutdown and the possible rise in the number of dynamic cases after it ends on May 3, 2020. Because the shutdown is still going on, our model predicts that the highest number of actively infected cases will be around 43,000 in the middle of May 2020. As a result of the government easing up on controls after the lockdown, we also expect the peak number of live tainted cases to rise by 7 to 21% in a variety of possible situations. India made the very

important choice to come up with a non-drug way to control the Coronavirus so that it doesn't spread during the busiest times and so that its general medical care system doesn't have to work too hard. For example, the whole country will be locked down for 40 days. As long as the Coronavirus spread continues, it is the responsibility of every country to come up with good public health and management strategies to fight the virus and protect their economies [5].

III. PROBLEM IDENTIFICATION

The identified problem in existing work is as follows:

(1) It may not be possible to identify COVID-19 cases because the accuracy is low.

(2) The COVID-19 patients' predictions may not be valid because they are hard to remember.

(3) If COVID-19 patient identification isn't done correctly, the accuracy may go down.

IV. RESEARCH OBJECTIVES

The research objectives as per identified problem in existing work are as follows:

(1) To improve precision for proper identification of COVID-19 patients.

(2) To improve recall for effective prediction of COVID-19 patients.

(3) To improve the accuracy of COVID-19 for exactness identification of patients.

V. METHODOLOGY

When a small sample is used to make a deep learning calculation, it often has an over-fitting problem. Our suggested plan is the best way to fix this problem and then work on making the CXR picture order more productive. Figure 4.4 shows the suggested method in the form of a block chart. DCRNN is a different kind of brain network that uses random noise to make fake picture ages by taking apart the parts of an info picture. It starts by focusing on the nearby highlights in the first few layers. After that, these local features get rid of the global ones.

Until the bottleneck, the encoder and decoder take turns down-sampling and up-sampling the data that is given. In the first four levels of the generator, the data is "down-examined," and then the features are learned. After the fifth layer, the picture is "up-tested," and so on. After that, the discriminator sorts the made picture into two groups: real pictures and engineering pictures. After the information is made, it is given as an input to the CNN sorting process.

Figure 1: Block Diagram of Proposed Method

Figure 2: Architecture of proposed CNN+DCRNN model.

Algorithm of Proposed Model

1. we need some input data; specifically, we need a picture as x and a label as y for our CXR actual photos. y = "typical," "pneumonia," and "COPD-IV-19."

2. Information provided as output Normal, pneumonia, or COVID-19 may be reported for output label y.

3. During the preprocessing stage, a 512 x 512 pixel resolution is applied to the CXR pictures.

For m steps of training, the number of iterations is as follows:

(i) Take a sample of m random numbers from the noise prior Pg(z);

(ii) take a sample of m random numbers from the distribution of generated examples Pdata(x).

The true picture is sent to the discriminator in step

(iii). Using the transfer model, the discriminator is updated by increasing its stochastic gradient.

conclusion for (i) The noise prior $Pg(z)$ is sampled in a minibatch of size m, and the discriminator is updated by minimizing its stochastic gradient.

The countdown has reached 5! In the testing step, a label y is produced as the output.

VI. RESULTS AND DISCUSSION

Python lets you set up deep learning projects where you can train networks in different ways at the start and then compare the results. Deep learning projects can be used to do things like

1. Go through a range of hyper-parameter values or use CNN to find the best ways to learn. For Bayesian optimization to work, you need the Statistics and Deep Learning Toolbox.

2. You can use the built-in function train network or make your own training function. If you use different data sets or try different classification network designs, you can compare the results.

3. If you use a template that has already been set up, you can get your project going quickly. Experiment templates can be used for workflows like image classification, image regression, sequence classification, semantic segmentation, and personalized training loops.

Filters can be used to narrow down and improve experiment results, and notes can be used to keep track of your findings. It's important that tests always give the same results. You can look at the descriptions of past studies to find the exact pairings of parameters that lead to each result.

Figure 5.1: Anaconda (Python 3.11) for proposed work

In above figure 3, shows the generalized code of the classification model which uses the concept of DCRNN-CNN.

Figure 4: Classification on x-ray data using proposed model

The above implementation shows that the classifies the train and test patient's data in different category.

The following observations are collected during the process of the proposed model on the patient X-Ray dataset. Precision, Recall, and Accuracy parameters are calculated as follows:

Table 1: Summary of results.

Figure 5: Graphical analysis of training and testing accuracy for COVID-19 X-Ray [1] dataset

The above graph show that the proposed model gives better training and testing accuracy as compare than AlexNet [1], GoogLeNet [1] and DCGAN-CNN [1] respectively. When use Covid X-Ray Dataset [1] then then training accuracy improve by 5.6%, 6.5%, 1.6% and testing accuracy improve by 4.9%, 8.7%, 2.1% for AlexNet [1], GoogLeNet [1] and DCGAN-CNN [1] respectively.

Figure 6: Graphical analysis of training and testing accuracy for COVID-19 Chest X-Ray [1] dataset

When use Covid Chest X-Ray Dataset [1] then then training accuracy improve by 4%, 1.6%, 1.6% and testing accuracy improve by 4%, 4.2%, 0.9% for AlexNet [1], GoogLeNet [1] and DCGAN-CNN [1] respectively.

When use Covid Radiography Dataset [1] then then training accuracy improve by 4%, 4.8%, 0.8% and testing accuracy improve by 3%, 4.3%, 1.1% for AlexNet [1], GoogLeNet [1] and DCGAN-CNN [1] respectively.

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Figure 8: Graphical analysis of train and test class accuracy in Normal cases for COVID-19 X-Ray [1] dataset

Figure 9: Graphical analysis of train and test class accuracy in Pneumonia cases for COVID-19 X-Ray [1] dataset

Figure 5.8: Graphical analysis of train and test class accuracy in Covid-19 cases for COVID-19 X-Ray [1] dataset

The above graph show that the proposed model gives better train and test class accuracy as compare than AlexNet [1], GoogLeNet [1] and DCGAN-CNN [1] respectively. When use Covid X-Ray Dataset [1] then train class accuracy improve by 6.7%, 5.9%, 1.1% and test class accuracy improve by 3.9%, 3.1%, 1.2% for AlexNet [1], GoogLeNet [1] and DCGAN-CNN [1] in Covid-19 cases.

Figure 10: Graphical analysis of train and test class accuracy in Normal cases for COVID-19 Chest X-Ray [1] dataset

Figure 11: Graphical analysis of train and test class accuracy in Pneumonia cases for COVID-19 Chest X-Ray [1] dataset

Figure 12: Graphical analysis of train and test class accuracy in Covid-19 cases for COVID-19 Chest X-Ray [1] dataset

When use Covid Chest X-Ray Dataset [1] then train class accuracy improve by 2.1%, 2.3%, 0.8% and test class accuracy improve by 1.6%, 4.3%, 0.85% for AlexNet [1], GoogLeNet [1] and DCGAN-CNN [1] in Covid-19 cases.

Figure 14: Graphical analysis of train and test class accuracy in Pneumonia cases for COVID-19 Radiography [1] dataset

Figure 15: Graphical analysis of train and test class accuracy in Covid-19 cases for COVID-19 Radiography [1] dataset

When use Covid Radiography Dataset [1] then train class accuracy improve by 1.8%, 5.2%, 0.7% and test accuracy improve by 2.5%, 6.2%, 0.5% for AlexNet [1], GoogLeNet [1] and DCGAN-CNN [1] in Covid-19 cases. Similarly, improvement of train and test class accuracy in normal and pneumonia cases.

Table 3: Recall and precision per class

Datas	Mod	Normal		Pneumon		COVID-	
et	el			ia		19	
		Re	Pre	Re	Pre	Re	Pre
		ca	cisi	ca	cisi	ca	cisi
		11	_{on}	11	on	11	on
COV	Alex	0.	0.8	0.	0.8	0.	0.9
ID-	Net	9	9	85	8	9	1
19 X-	$[1]$						
ray							
$[1]$	Goo	0.	0.9	0.	0.8	$\overline{0}$.	0.8
	gLe	91		86	9	87	8
	Net						
	$[1]$						
	DCG	$\overline{0}$.	0.9	$\boldsymbol{0}$.	0.8	$\overline{0}$.	0.9
	AN-	92	3	93	9	95	\overline{c}
	CNN						
	$[1]$						
	Prop	$\overline{0}$.	0.9	0.	0.9	$\overline{0}$.	0.9
	osed	94	5	97	\overline{c}	96	$\overline{4}$
	DCR						
	NN-						
	CNN						

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Figure 16: Graphical analysis of recall in Normal, Pneumonia and Covid-19 cases for COVID-19 X-Ray [1] dataset

Figure 17: Graphical analysis of precision in Normal, Pneumonia and Covid-19 cases for COVID-19 X-Ray [1] dataset

The above graph show that the proposed model gives better recall and precision as compare than AlexNet [1], GoogLeNet [1] and DCGAN-CNN [1] respectively. When use Covid X-Ray Dataset [1] then recall improve by 6.7%, 10.3%, 1.05% and precision improve by 3.3%, 6.8%, 2.2% for AlexNet [1], GoogLeNet [1] and DCGAN-CNN [1] in Covid-19 cases.

Figure 18: Graphical analysis of recall in Normal, Pneumonia and Covid-19 cases for COVID-19 Chest X-Ray [1] dataset

Figure 19: Graphical analysis of precision in Normal, Pneumonia and Covid-19 cases for COVID-19 Chest X-Ray [1] dataset

When use Covid Chest X-Ray Dataset [1] then recall improve by 3.2%, 8.7%, 2.2% and precision improve by 6.7%, 5.5%, 2.1% for AlexNet [1], GoogLeNet [1] and DCGAN-CNN [1] in Covid-19 cases.

Figure 20: Graphical analysis of recall in Normal, Pneumonia and Covid-19 cases for COVID-19 Radiography [1] dataset

Figure 2: G1raphical analysis of precision in Normal, Pneumonia and Covid-19 cases for COVID-19 Radiography [1] dataset

When use Covid Radiography Dataset [1] then recall improve by 4.3%, 8.9%, 2.1% and precision improve by 3.2%, 6.5%, 1.04% for AlexNet [1], GoogLeNet [1] and DCGAN-CNN [1] in Covid-19 cases. Similarly, improvement of recall and precision in normal and pneumonia cases.

VII. CONCLUSIONS

These are supposed to have conclusions, and this one do:

(1) AlexNet [1], GoogLeNet [1], and DCGAN-CNN [1] all saw increases in training accuracy of 5.6%, 6.5%, and 1.6%, and in testing accuracy of 4.9%, 8.7%, and 2.1%, respectively.

(2) For Covid-19, AlexNet [1], GoogLeNet [1], and DCGAN-CNN [1] each show gains of 6.7%, 5.9%, 1.1% in train class accuracy and 3.9%, 3.1%, 1.2% in test class accuracy, respectively.

(3) The proposed model outperforms AlexNet [1], GoogLeNet [1], and DCGAN-CNN [1] in terms of recall and accuracy. AlexNet [1], GoogLeNet [1], and DCGAN-CNN [1] all perform better on the Covid X-Ray Dataset [1], with recall increasing by 6.7%, 10.3%, and 1.05% and precision increasing by 3.3%, 6.8%, and 2.2% in the Covid-19 instances, respectively.

Recall and accuracy have also increased in both nonpneumonia patients and those with pneumonia. Therefore, the suggested system, DCRNN-CNN (Deep Convolution Recurrent Neural Network with CNN), is more effective in classifying X-Ray pictures for normal, pneumonia, and Covid-19 illness symptoms.

The diagnostic precision and treatment efficacy are both enhanced by the approach we offer. Future improvements must evaluate accuracy with new datasets and use other AI methods to verify precision calculations. Due to the massive quantity of data needed for performance estimation of train data, the proposed model has a processing time restriction. In the future, the same algorithms will be used to data in real time to estimate the system's efficacy.

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