

COVID-19 CHEST CT IMAGE SEGMENTATION NETWORK BY MULTI SCALE FUSION AND ENHANCEMENT OPERATIONS

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ABSTRACT-A deep learning research is the availability of training data sets. With a limited number of publicly available COVID-19 chest X-ray images, the generalization and robustness of deep learning models to detect COVID-19 cases developed based on these images are questionable. Segmentation of pneumonia lesions from Lung CT images has become vital for diagnosing the disease and evaluating the severity of the patients during the COVID-19 pandemic. Several Deep Learning based systems have been proposed for this task. However, some low-contrast abnormal zones in CT images make the task challenging. which will be used as the testing data set. We used a deep learning model based on the convolutional neural network architecture, which was pre trained to recognize objects from a million of images and then retrained to detect abnormality in chest X-ray images.

Keywords—Deep Learning, COVID-19, pneumonia prediction, Evolution, Intelligent System.

1.INTRODUCTION

The World Health Organization (WHO) has declared the corona virus disease 2019 (COVID-19)[1] a pandemic. A global coordinated effort is needed to stop the further spread of the virus. A pandemic is defined as “occurring over a wide geographic area and affecting an exceptionally high proportion of the population. The last pandemic reported in the world was the H1N1 flu pandemic in 2009. On 31 December 2019, a cluster of cases of pneumonia of unknown cause, in the city of Wuhan, Hubei province in China, was reported to the World Health Organization. In January 2020, a previously unknown new virus was identified, subsequently named the 2019 novel corona virus, and samples obtained from cases and analysis of the virus’ genetics indicated that this was the cause of the outbreak. This novel corona virus was named Corona virus Disease 2019 (COVID-19) by WHO in February 2020. The virus is referred to as SARS-CoV-2 and the associated disease is COVID.

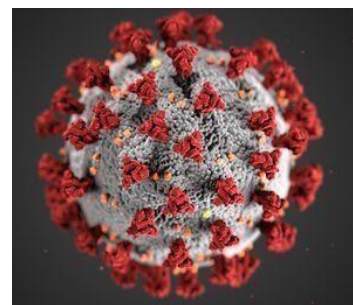


Fig. 1.1 Corona Virus

Corona viruses are a family of viruses that cause illness such as respiratory diseases or gastrointestinal diseases. Respiratory diseases can range from the common cold to more severe diseases. Example:

- 1) Middle East Respiratory Syndro
- 2) Acute Respiratory Syndrome (SARS-CoV).

A novel corona virus (nCoV) is a new strain that has not been identified in humans previously. Once scientists determine exactly what corona virus it is, they give it a name (as in the case of COVID-19, the virus causing it is SARS-CoV- 2). Corona viruses got their name from the way that they look under a microscope. The virus consists of a core of genetic material surrounded by an envelope with protein spikes. This gives it the appearance of a crown. The word Corona means “crown” in Latin. Corona viruses are zoometric, meaning that the viruses are transmitted between animals and humans. It has been determined that MERS-CoV was transmitted from dromedary camels to humans and SARS-CoV from civet cats to humans. The source of the SARS-CoV-2 (COVID-19) is yet to be determined, but investigations are ongoing to identify the zoonotic source to the outbreak.

WHO advice for high-risk populations:

- When having visitors at your home, extend “1-meter greetings”, like a wave, nod or bow.
- Request that visitors and those who live with you, wash their hands.
- Clean and disinfect surfaces in your home (especially those that people touch a lot) on a regular basis.
- Limit shared spaces if someone you live with is not feeling well (especially with possible COVID-19 symptoms).
- If you show signs and symptoms of COVID-19 illness, contact your healthcare provider by telephone, before visiting your healthcare facility.
- Have an action plan in preparation for an outbreak of COVID-19 in your community.
- When you are in public, practice the same preventative guidelines as you would at home.
- Keep updated on COVID-19 through obtaining information from reliable sources.

Transmission of COVID-19

Evidence is still emerging, but current information is indicating that human-to-human transmission is occurring. The routes of transmission of COVID-19 remains unclear at present, but evidence from other corona viruses and respiratory diseases indicates that the disease may spread through large respiratory droplets and direct or indirect contact with infected secretions.

The incubation period of COVID-19 is currently understood to be between 2 to 14 days. This means that if a person remains well after 14 days after being in contact with a person with confirmed COVID-19, they are not infected.

Literature review (June 2020) investigates and discusses the unclear issues related to disease transmission and pathogenesis and the accuracy of diagnostic tests and treatment modalities.

Differential Diagnosis

Differential diagnosis should include the possibility of a wide range of common respiratory disorders such as: Other Corona viruses

- Respiratory
- Syncyt
- Rhinovirus
- Bacterial pneumonia, mycoplasma pneumonia (MPP)
- Chlamydia pneumonia.

Differentiation should also be made from lung disease caused by other diseases. A CT scan has great value in early screening and differential diagnosis [2] for COVID-19.

Management / Interventions

In the case of mild to moderate symptoms the following considerations should be taken into account:

Early identification - Clinicians, especially physiotherapists, are most often in direct contact with their patients, which can make them infected or infected by others. It is therefore very important for physiotherapists and other health professionals to be familiar with the condition of COVID-19, how to identify it and how to prevent it.

strategies for infection prevention and control (IPC) - Suspect, probable and confirmed cases should be educated on IPC strategies to prevent transmission of the disease and health management strategies for quarantine.

Diagnostic Procedures

A COVID-19 diagnostic testing kit has been developed and is available in clinical testing labs. The gold standard for testing for COVID-19 is Reverse Transcription Polymerase Chain Reaction (RT-PCR). However, current data suggest that RT-PCR is only 30-70% effective for acute infection, this may be due to incorrect use of lab kits or not enough virus in the blood at the early stages of testing. Plus, the availability of testing will vary from country to country. The CDC recommends that any person who may have had contact with a person who is suspected of having COVID-19 and develops a fever and respiratory symptoms listed above are advised to call their healthcare practitioner to determine the best course of action.

If the above criteria are met it is advised that the following testing procedure is followed: Collect and test upper respiratory tract specimens, using a nasopharyngeal swab. If available testing of lower respiratory tract specimens. If a productive cough is evident then a sputum specimen should be collected. For patients who are receiving invasive mechanical ventilation, a lower respiratory tract aspirate or broncho-alveolar lavage sample should be collected.

II. LITERATURE SURVEY

1. “Marginal Space Deep Learning: Efficient Architecture for Volumetric Image Parsing”

Florin C. et al proposed a pipeline for object detection and segmentation in the context of volumetric image parsing, solving a two-step learning problem: anatomical pose estimation and boundary delineation. For this task we introduce Marginal Space Deep Learning (MSDL), a novel framework exploiting both the strengths of efficient object parametrization in hierarchical marginal spaces and the automated feature design of Deep Learning (DL) network architectures. In the 3D context, the application of deep learning systems is limited by the very high complexity of the parametrization. More specifically 9 parameters are necessary to describe a restricted affine transformation in 3D, resulting in a prohibitive amount of billions of scanning hypotheses. To further increase computational efficiency and robustness, in our system we learn sparse adaptive data sampling patterns that automatically capture the structure of

the input. Given the object localization, we propose a DL-based active shape model to estimate the non-rigid object boundary. Experimental results are presented on the aortic valve in ultrasound using an extensive dataset of 2891 volumes from 869 patients, showing significant improvements of up to 45.2%

2. “Deep Learning Based Detection and Correction of Cardiac MR Motion Artefacts During Reconstruction for High-Quality Segmentation”

Ilkay Oksuz et al discuss the implications of image motion artefacts on cardiac MR segmentation and compare a variety of approaches for jointly correcting for artefacts and segmenting the cardiac cavity. The method is based on our recently developed joint artefact detection and reconstruction method, which reconstructs high quality MR images from k-space using a joint loss function and essentially converts the artefact correction task to an under-sampled image reconstruction task by enforcing a data consistency term. In this paper, we propose to use a segmentation network coupled with this in an end-to-end framework. Our training optimises three different tasks: 1) image artefact detection, 2) artefact correction and 3) image segmentation. We train the reconstruction network to automatically correct for motion-related artefacts using synthetically corrupted cardiac MR k-space data and uncorrected reconstructed images. Using a test set of 500 2D+time cine MR acquisitions from the UK Biobank data set, we achieve demonstrably good image quality and high segmentation accuracy in the presence of synthetic motion artefacts. We showcase better performance compared to various image correction architectures.

3. “Deep Learning for Classification and Localization of COVID-19 Markers in Point-of-Care Lung Ultrasound”

Subhankar Roy et al studies the application of DL techniques for the analysis of lung ultrasonography (LUS) images. Specifically, we present a novel fully-annotated dataset of LUS images collected from several Italian hospitals, with labels indicating the degree of disease severity at a frame-level, video-level, and pixel-level (segmentation masks). Leveraging these data, we introduce several deep models that address relevant tasks for the automatic analysis of LUS images. In particular, we present a novel deep network, derived from Spatial Transformer Networks, which simultaneously predicts the disease severity score associated to a input frame and provides localization of pathological artefacts in a weakly-supervised way. Furthermore, we introduce a new method based on uninform for effective frame score aggregation at a video-level. Finally, we benchmark state of the art deep models for estimating pixel-level segmentations of COVID-19 imaging biomarkers. Experiments on the proposed dataset demonstrate satisfactory results on all the considered tasks, paving the way to future research on DL for the assisted diagnosis of COVID-19 from LUS data. However, application of these models in clinically realistic environments can result in poor generalization and decreased accuracy, mainly due to the domain shift across different hospitals, scanner vendors, imaging protocols, and patient populations etc.

4. “Automatic 3D Bi-Ventricular Segmentation of Cardiac Images by a Shape-Refined Multi- Task Deep Learning Approach”

Jinming Duan et al proposed a multi-task deep learning approach with atlas propagation to develop a shape-refined bi-ventricular segmentation pipeline for short-axis CMR volumetric images. The pipeline first employs a fully convolutional network (FCN) that learns segmentation and landmark localization tasks simultaneously. The architecture of the proposed FCN uses a 2.5D representation, thus combining the computational advantage of 2D FCNs networks and the capability of addressing 3D spatial consistency without compromising segmentation accuracy. Moreover, a refinement step is designed to explicitly impose shape prior knowledge and improve segmentation quality. This step is effective for overcoming image artifacts (e.g., due to different breath-hold positions and large slice thickness), which preclude the creation of anatomically meaningful 3D cardiac shapes. The pipeline is fully automated, due to network's ability to infer landmarks, which are then used downstream in the pipeline to initialize atlas propagation. We validate the pipeline on 1831 healthy subjects and 649 subjects with pulmonary hypertension. Extensive numerical experiments on the two datasets demonstrate that our proposed method is robust and capable of producing accurate, high-resolution, and anatomically smooth bi-ventricular 3D models, despite the presence of artifacts in input CMR volumes.

5. “Generalizing Deep Learning for Medical Image Segmentation to Unseen Domains via Deep Stacked Transformation”

Specifically, a series of n stacked transformations are applied to each image during network training. The underlying assumption is that the “expected” domain shift for a specific medical imaging modality could be simulated by applying extensive data augmentation on a single source domain, and consequently, a deep model trained on the augmented “big” data (BigAug) could generalize well on unseen domains. We exploit four surprisingly effective, but previously understudied, image-based characteristics for data augmentation to overcome the domain generalization problem. We train and evaluate the BigAug model (with $n=9$ transformations) on three different 3D segmentation tasks (prostate gland, left atrial, left ventricle) covering two medical imaging modalities (MRI and ultrasound) involving eight. The underlying assumption is that the “expected” domain shift for a specific medical imaging modality could be simulated by applying extensive data augmentation on a single source domain, and consequently, a deep model trained on the augmented “big” data (Big Aug) could generalize well on unseen domains. We exploit four surprisingly effective, but previously understudied, image-based characteristics for data augmentation to overcome the domain generalization problem.

6. “Deep Neural Network Regression for Automated Retinal Layer Segmentation in Optical Coherence Tomography Images”

Lua Ngo et al propose an automated segmentation method for OCT images based on a feature-learning regression network without human bias. The proposed deep neural network regression takes the intensity, gradient, and adaptive normalized intensity score (ANIS) of an image segment as features for learning, and then predicts the corresponding retinal boundary pixel. Reformulating the segmentation as a regression problem obviates the need for a huge dataset and reduces the complexity significantly, as shown in the analysis of computational complexity given here. In addition, assisted by ANIS, the method operates robustly on OCT images containing intensity variances, low-contrast regions, speckle noise, and blood vessels, yet remains accurate and time-efficient. In the evaluation of the method conducted using 114 8. “Parallel Deep Learning Algorithms with Hybrid Attention Mechanism for Image Segmentation of Lung Tumors”

Hexuan Hu et al proposed a parallel deep learning algorithm with hybrid attention mechanism for image segmentation. Firstly, lung parenchyma was extracted via preprocessing images. Then, images were input into hybrid attention mechanism and DenseNet module, respectively, where hybrid attention mechanism consisted of a spatial attention mechanism and a channel attention mechanism. Finally, four feasible solutions were proposed for the verification through changing the convolution quantity of Dense block in DenseNet. The network structure with the better performance was achieved. The experimental results prove the parallel deep learning algorithm with hybrid attention mechanism performed well in image segmentation of lung tumors, and its accuracy can reach 94.61%.

III. PROPOSED METHODOLOGY

In this project we present an evaluation of pre-trained CNN model in a transfer learning setup for COVID-19 detection from chest X-ray images. System should be Self sustained without Doctor Intervention. Should have More accuracy and Precision.

The segmented primary and secondary regions are compressed with hybrid techniques for telemedicine application.

Lung Nodule Detection

A lung nodule (or mass) is a small abnormal area that is sometimes found during a CT scan of the chest. These scans are done for many reasons, such as part of lung cancer screening, or to check the lungs if you have symptoms.

Pneumonia Detection

A cough is caused by irritation in the nerves in your airway. A chronic cough, which may require a chest x-ray, lasts 8 weeks or more. Sometimes a persistent cough can be an indication of conditions like pneumonia, which can be diagnosed through a chest x-ray.

size, all images need to be resized to a pixel size before inputting them to the CNN.

Feature Extraction:

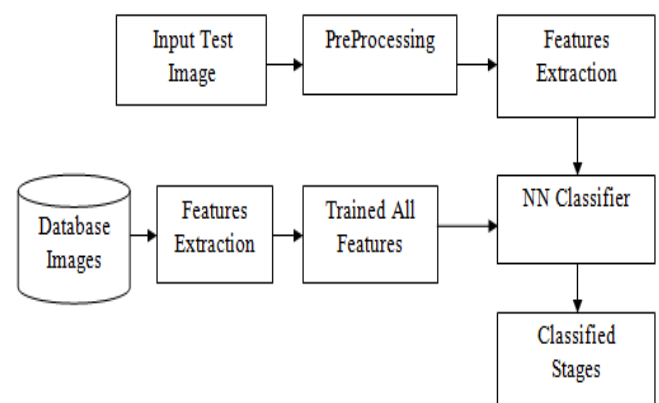
Next thing is to do Feature extraction is an attribute reduction process. Unlike feature selection, which ranks the existing attributes according to their predictive significance, feature extraction actually transforms the attributes. The transformed attributes, or features, are linear combinations of the original attributes. Finally, our models are trained using Classifier algorithm. We use classify module on Natural Language Toolkit library on Python. We use the labeled dataset gathered. The rest of our labeled data will be used to evaluate the models. Some machine learning algorithms were used to classify pre-processed data.

The technique of extracting the features is useful when you have a large data set and need to reduce the number of resources without losing any important or relevant information.

Classification

There is a fluctuation in the performance of these techniques as well depending on the data and other prerequisite steps. However, in this research work, classification is performed through CNN. This algorithm is referred to as a supervised machine learning approach that is commonly used for classification and regression problems. CNN is an efficient algorithm that shows encouraging results on a given dataset.

BLOCK DIAGRAM



DEEP LEARNING

A small subset of Artificial Intelligence (AI), often called Machine Learning (ML)[5], has revolutionized several fields in the last few decades. Neural Networks (NN[6]) is a subfield of ML, and it was this subfield that spawned Deep Learning (DL). Since its inception DL has been creating ever larger disruptions, showing outstanding success in almost every application domain. Figure 1 shows the taxonomy of AI. DL which uses either deep architecture of learning or hierarchical learning approaches, is a class of ML developed largely from 2006 onward. Learning is a procedure consisting of estimating the model parameters so that the learned model (algorithm) can perform a specific task. For example, in Artificial Neural Networks (ANN), the parameters are the weight matrices. DL, on the other hand, consists of several layers in between the input and output layer which allows for many stages of non-linear information processing units with hierarchical architectures to be present that are exploited for feature learning and pattern classification.

Learning methods based on representations of data can also be defined as representation learning. Recent literature states that DL based representation learning involves a hierarchy of features or concepts, where the high-level concepts can be defined from the low-level ones and low-level concepts can be defined from high-level ones. In some articles, DL has been described as a universal learning approach that is able to solve almost all kinds of problems in different application domains. In other words, DL is not task specific.

Type of Deep Learning Approaches

Deep learning approaches can be categorized as follows: Supervised, semi-supervised or partially supervised, and unsupervised. In addition, there is another category of learning approach called Reinforcement Learning (RL) or Deep RL (DRL) which are often discussed under the scope of semi-supervised or sometimes under unsupervised learning approaches. Fig 1.3 shows the pictorial diagram. AI: Artificial Intelligence; ML: Machine Learning; NN: Neural Networks; DL: Deep Learning; SNN: Spiking Neural Networks

Deep Supervised Learning

Supervised learning is a learning technique that uses labeled data. In the case of supervised DL approaches, the environment has a set of inputs and corresponding outputs $(x_t, y_t) \sim p$. For example, if for input x_t , the intelligent agent predicts $y_t = f(x_t)$, the agent will receive a loss value $l(y_t, y^*_t)$. The agent will then iteratively modify the network parameters for a better approximation of the desired outputs. After successful training, the agent will be able to get the correct answers to questions from the environment.

There are different supervised learning approaches for deep learning, including Deep Neural Networks (DNN), Convolutional Neural Networks (CNN), Recurrent Neural Networks (RNN), including Long Short Term Memory (LSTM), and Gated Recurrent Units (GRU). These networks will be described in details in the respective sections.

Deep Semi-supervised Learning

Semi-supervised learning is learning that occurs based on partially labeled datasets. In some cases, DRL and Generative Adversarial Networks (GAN)[7] are used as semi-supervised learning techniques.

Deep Unsupervised Learning

Unsupervised learning systems are ones that can without the presence of data labels. In this case, the agent learns the internal representation or important features to discover unknown relationships or structure within the input data. Often clustering, dimensionality reduction, and generative techniques are considered as unsupervised learning approaches. There are several members of the deep learning family that are good at clustering and non-linear dimensionality reduction, including Auto-Encoders (AE), Restricted Boltzmann Machines (RBM), and the recently developed GAN. In addition, RNNs, such as LSTM and RRL, are also used for unsupervised learning in many application domains.

Deep Reinforcement Learning (RL)

Deep Reinforcement Learning is a learning technique for use in unknown environments. DRL began in 2013 with Google Deep Mind. From then on, several advanced methods have been proposed based on RL. Here is an example of RL: If environment samples inputs $x_t \sim p$, agent predicts $y_t = f(x_t)$, agent receive cost: $c_t \sim P(c_t | x_t, y_t)$ where P is an unknown probability distribution, the environment asks an agent a question, and gives a noisy score as the answer. Sometimes this approach is called semi-supervised learning as well. There are many semi-supervised and un-supervised techniques that have been implemented based on this concept. In RL, we do not have a straight forward loss function, thus making learning harder compared to traditional supervised approaches. The fundamental differences between RL and supervised learning are: First, you do not have full access to the function you are trying to optimize; you must query them through interaction, and second, you are interacting with a state-based environment: Input x_t depends on previous actions. Depending upon the problem scope or space, one can decide which type of RL needs to be applied for solving a task. If the problem has a lot of parameters to be optimized,

DRL is the best way to go. If the problem has fewer parameters for optimization, a derivation free RL approach is good. An example of this is annealing, cross entropy methods, and SPSA.

CNN OVERVIEW-CNN: In deep learning, a convolution neural network (CNN/Convent)[8] is a class of deep neural networks, most commonly applied to analyze visual imagery. Now when we think of a neural network we think about matrix multiplications but that is not the case with Convent. It uses a special technique called Convolution. Now in mathematics convolution is a mathematical operation on two functions that produces a third function that expresses how the shape of one is modified by the other. Background of CNNs- CNN's were first developed and used around the 1980s. The most that a CNN[9] could do at that time was recognizing handwritten digits. It was mostly used in the postal sectors to read zip codes, pin codes, etc. The important thing to remember about any deep learning model is that it requires a large amount of data to train and also requires a lot of computing resources. This was a major drawback for CNNs at that period and hence CNNs were only limited to the postal sectors and it failed to enter the world of machine learning **SVM-A** support vector machine (SVM) is a type of supervised learning algorithm used in machine learning to solve classification and regression tasks; SVMs are particularly good at solving binary classification problems, which require classifying the elements of a data set into two groups.

Network Parameters and Required Memory for CNN

The number of computational parameters is an important metric to measure the complexity of a deep learning model. The size of the output feature maps can be formulated as follows:

$$M = \frac{(N - F)}{S} + 1,$$

where N refers to the dimensions of the input feature maps, F refers to the dimensions of the filters or the receptive field, M refers to the dimensions of output feature maps, and S stands for the stride length. Padding is typically applied during the convolution operations to ensure the input and output feature map have the same dimensions. The amount of padding depends on the size of the kernel'shis Equation is used for determining the number of rows and columns for padding.

$$P = (F - 1)/2,$$

here P is the amount of padding and F refers to the dimension of the kernels. Several criteria are considered for comparing the models. However, in most of the cases, the number of network parameters and the total amount of memory are considered. The number of parameters (Pram) of lthlayer is the calculated basedon the following equation:

$$\begin{aligned} \text{Parm}_i &= (F * F * \text{FM}_{i-1}) * \text{FM}_i \\ \text{If bias is added with the weights, then the above} \\ \text{equation can be written as follows:} \\ \text{Parm}_i &= (F * (F + 1) * \text{FM}_{i-1}) * \text{FM}_i \end{aligned}$$

Residual Network (Reset in 2015)

The winner of ILSVRC 2015 was the Residual Network[9] architecture, ResNet. Resnet was developed by Kaiming He with the intent of designing ultra- deep networks that did not suffer from the vanishing gradient problem that predecessors had. ResNet is developed with many different numbers of layers; 34, 50, 101, 152, and even 1202. The popular ResNet50 contained 49 convolution layers and 1 fully connected layer at the end of the network. The total number of weights and MACs for the whole network is 25.5M and 3.9M respectively. The basic block diagram of the ResNet architecture is shown in Figure 16. ResNet is a traditional feed forward network with a residual connection. The output of a residual layer can be defined based on the outputs of (l- 1) the which comes from the previous layer defined as x_{l-1} . $\mathcal{F}(x_{l-1})$ is the output after performing various operations (e.g., convolution with different size of filters, Batch Normalization (BN) followed by an activation function, such as a Rule on x_{l1}). The final output of unit is

x_l which can be defined with the following equation: $x_l = \mathcal{F}(x_{l-1}) + x_{l-1}$. (15)

The residual network consists of several basic residual blocks. However, the operations in the residual block can be varied depending on the different architecture of residual networks. The wider version of the residual network was proposed by Zagoruvko et al., another improved residual network approach known as aggregated residual transformation. Recently, some other variants of residual models have been introduced based on the Residual Network

architecture. Furthermore, there are several advanced architectures that are combined with Inception and Residual units. The basic conceptual diagram of Inception-Residual unit is shown in the following.

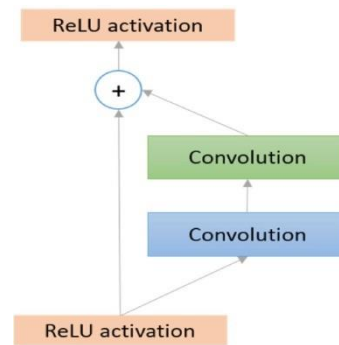
IV.RESULT & DISCUSSION

We employed the simulation on the Kaggle Covid-19 dataset for machine learning and deep learning methods.^{1,3} The experiment mainly finds out the various features include texture, morphological, and image quality to distinguish Covid-19 infected patients and normal person based on lung infections. The main finding of the research was able to detect the Covid-19 infected patients among two popular Machine learning (Random forest) and Deep learning (CNN) methods.

In the previous research^{3,4} machine learning methods and research paper^{1,2} deep learning methods were used. It is crucial to formulate a deep neural network system like COVID-Net with transparency and accountability in view, because of the mission-critical nature of biomedical studies such as COVID-19 identification that really can affect patients' wellness and well-being. This research work evaluated the performance of machine learning-based random forest and deep learning-based CNN method. In the existing research²⁻⁴ researchers are mainly utilizing the machine learning method, which performs better when the data size is limited. The precision of the experiment, together with the complexity of the design, the number of parameters, and the complexity of data processing are shown in Tables 2 and 3. This can be noticed that just by accomplishing more than 90 % test accuracy of CNN achieves better accuracy, consequently illustrating the effectiveness of exploiting participatory management framework strategies to build fully personalized deep neural network system in an increased way made to fit assignment, statistics, and strategic goals.

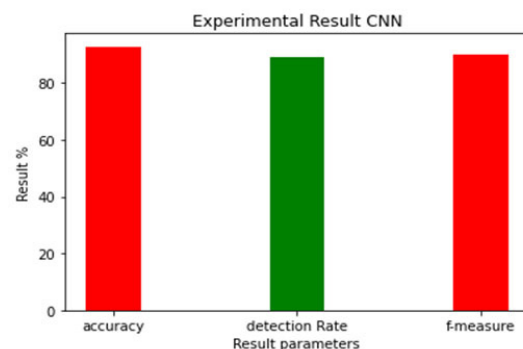
The various experimental parameters include detection rate, accuracy, and F-measure were calculated for CNN (Deep learning) and Random Forest (Machine learning method). The leading multi-class characteristic was the boundary. This same boundary seems to be the total pixel quantity at the edge of a picture. The fastest-growing functionalities from COVID-19 Vs viral infection were skewness, randomness, compact size, and thin proportion. The experimental results The CNN model outperforms over Random forest method, it shows an accuracy of 95.4 %

Figures 2-5) by considering the 40 % of training data and 60 % of testing data for the Covid-19 X-ray data set. Together, these results indicate that the boundary is a key differentiated characteristic, coherent with a closely examining which lung infection with COVID-19 appears to become more extraneous and lateral along the lung borders.



To determine a better understanding of the overlap between all the decision-making procedure of deep neural networks [10] besides potential treatments as well as the policy procedure of healthcare professionals throughout chest radiographs, these observations are therefore informative. All performance indicators fell substantially although expected mostly with multi-class classification. Even so, the Classifier and accuracy coupled remained high. These research results are enabling and recommend that COVID-19 lung infection can indeed be distinguished from many other comparable lung infections by the multi-class categorization.

The purpose of this research was to enhance forecasting power by retrieving texture as well as morphological characteristics from X-ray images. The process of machine-learning is a hard process to retrieve the investigators' most suitable and important characteristics. Findings demonstrate that the most suitable and important hidden details provided in the COVID-19 lung infection, which also enhanced the multiple as well as inter classification, is available throughout the segmented images utilizing the CNN approach. These characteristics are most often used as feedback to the CNN process. The findings acquired outperformed these earlier conventional techniques.



V.CONCLUSION & FUTURE WORK

The outbreaks of COVID-19 is not even the first pandemic, as well as doubtful to be last. However, over the first time,

modern communities get the resources that provide a structured, evidence-based, equitable, as well as international solution towards human health. If we try to analyze these images manually it takes lots of time as well as some possibility of inaccuracy. Due to the rapid growth of the Covid-19 pandemic and limited resources (detection kits, lab, health workers, and medicines), the need for an automatic and accurate detection system is always an interesting hot topic for researchers. By using the effectiveness of the whole solution increasing depends partially on ML and DL methods. This research mainly utilizes various chest images (X-ray) of covid-19 patients. The experimental results influence that the CNN method shows a 92.4 % accuracy result over the machine learning method. We could also save numerous human lives now and in the future, unless we begin taking the whole chance to acquire data, pool our expertise as well as incorporate our skill and knowledge. Throughout future work, we will build a methodology premised on artificial intelligence and machine learning to battle in monitoring and mitigation with pandemic COVID-19.

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