



PNEUMONIA PREDICTION AND DETECTION USING MACHINE LEARNING ALGORITHM

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ABSTRACT:

Pneumonia is a respiratory disease caused by bacteria or viruses. It affects many people, especially in developing and developing countries, where pollution levels are high, unhealthy living conditions and congestion are common along with inadequate medical infrastructure. Pneumonia causes pleural leakage, a condition in which fluid leaks out fill the lungs, causing difficulty breathing. Early diagnosis of pneumonia is important to confirm therapeutic treatment and increasing survival rates. Chest X-ray imaging is the most common method used to diagnose pneumonia. However, chest X-ray examination is a challenging work and prone to tangible flexibility.

Pneumonia remains a threat to human health; Coronavirus 2019 (COVID-19) which started in late 2019 has had a huge impact on the world. It continues in many lands and has caused tremendous losses in human life and property. In this paper, we present the DeepConv-DilatedNet-based method of diagnosis and locating pneumonia in chest X-ray images (CXR).

The two-phase Faster R-CNN detector was adopted as a network infrastructure. Feature Pyramid Network (FPN) is integrated into an expanded neural network of bottleneck extensions to deepen features to preserve in-depth feature and location information. In the case of DeepConv-DilatedNet, the deconvolution network is used to restore high-level feature maps to their original size, and targeted information is stored.

DeepConv-DilatedNet, on the other hand, uses the most popular full-featured layouts and shares some calculations throughout the image. Then, Soft-NMS is used to check boxes and verify sample quality. Also, K-Means ++ is used to produce work boxes to improve the precision of spatial processing. The algorithm obtained 39.23% Mean Average Precision (mAP) from the X-ray image database from the Radiological Society of North America (RSNA) and obtained 38.02% Mean Average Precision (mAP) from the ChestX-ray14 database, surpassing other information algorithms. Therefore, in this paper, an advanced algorithm that can provide doctors with information on the location of pneumonia lesions is proposed.

KEYWORDS: Pneumonia, Machine Learning, deep LEARNING, DeepConv-DilatedNet-based method, R-CNN detector, Soft-NMS, K-Means ++ algorithm, CXR, mAP.

1. INTRODUCTION:

Machine learning (ML) is a field of inquiry devoted to understanding and building methods that 'learn', that is, methods that leverage data to improve performance on some set of tasks.[1] It is seen as a part of artificial intelligence. Machine learning algorithms build a model based on sample data, known as training data, in order to make predictions or decisions without being explicitly programmed to do so.[2] Machine learning algorithms are used in a wide variety of applications, such as in medicine, email filtering, prediction, diagnosis, speech recognition, and computer vision, where it is difficult or unfeasible to develop conventional algorithms to perform the needed tasks. Over the past two decades Machine Learning has become one of the main- stays of information technology and with that, a rather central, albeit usually hidden, part of our life. With the ever increasing amounts of data becoming available there is good reason to believe that smart data analysis will become even more pervasive as a necessary ingredient for technological progress.

The purpose of this paper is to understand the machine learning applications in medical diagnosis especially in pneumonia prediction and detection. For target detection, first, we chose the two-stage model Faster R-CNN with a suitable detection effect as the prediction framework, and then, we considered the excellent effect of DetNet59 as the backbone, so we used DetNet59 for the feature extraction of the target. But because DetNet59 up sampling will lose features, we improve DetNet59 and change up sampling to deconvolution to reduce feature loss and at the same time add whole convolution to expand the receptive field. Besides, the initial anchor box has a great influence on the training and prediction of the model, so we borrowed from the YOLO method and used the K-Means++ algorithm to obtain the initial anchor box for the pneumonia target dataset to improve the detection effect. Finally, because the Soft-NMS improves the performance of the target detection network, we have joined Soft-NMS.

Pneumonia is a dangerous lung disease that can be caused by germs, germs, or fungi and enters the lungs, causing inflammation of the air sacs and pleural effusion, a condition when the lungs are filled with fluid. It causes more than 15% of child deaths in less than five years [1]. Pneumonia is most common in less developed and developed countries, where overcrowding, pollution, and unsanitary conditions. Exacerbate the situation and medical resources are limited. Therefore, the diagnosis is still early and management can play an important role in preventing the disease from killing. Radiological examination of the lungs using computed tomography (CT), magnetic resonance imaging (MRI), or radiography (X-ray) is often used for diagnosis. X-ray imaging creates a less aggressive and less expensive lung test. figure 1 shows an example of an X-ray of the lungs and a healthy lung. White dots on X-ray radiation (indicated by red arrows), called infiltrates, separate pneumonia healthy state. b shows the X-ray of a pneumonia affected lung.

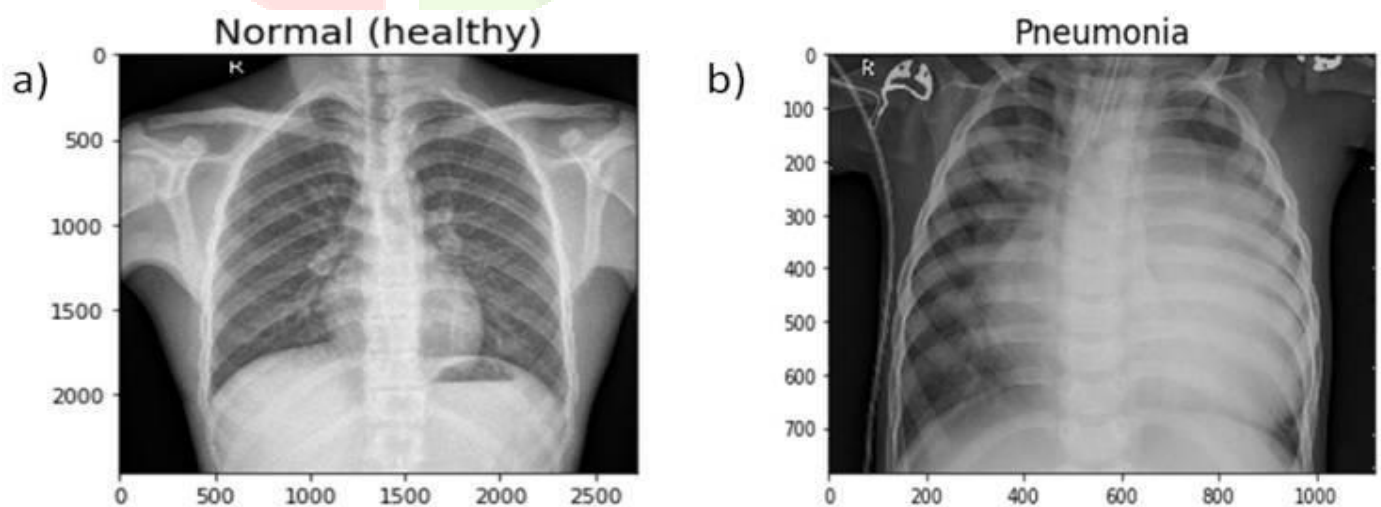


Figure 1 Examples of two X-ray plates show (a) healthy lung and (b) pneumonia. The red arrows in (b) indicate the white entry, the distinguishing feature of pneumonia.

Table 1

Description of the experimental dataset.

Category	Training Set	Test Set	11.95%
Normal (Healthy)	1283	300	
Pneumonia (Viral + Bacteria)	3873	400	
Total	5156	700	
			05%

The dataset comprised a total of 5836 images (table1) segmented into two main parts, a training set and a test set. Both bacterial and viral pneumonia were considered as a single category, pneumonia infected.

The dataset used in this study did not include any case of viral and bacterial co-infection. All chest X-ray images were taken during the routine clinical care of the patients. Two expert physicians then graded the diagnoses for the images before being cleared for training the AI system. The evaluation set was also checked by a third expert to account for any grading errors. The proportion of data assigned to training and testing was highly imbalanced. Therefore, the dataset was shuffled and arranged into training and test sets. Finally, there were 5136 images in the training set and 700 images in the test set and then Eleven-point-nine-five percent of the complete dataset was used as the testing dataset. Figure 1 shows two chest X-ray images, one of a healthy person and the other of a person suffering from pneumonia.

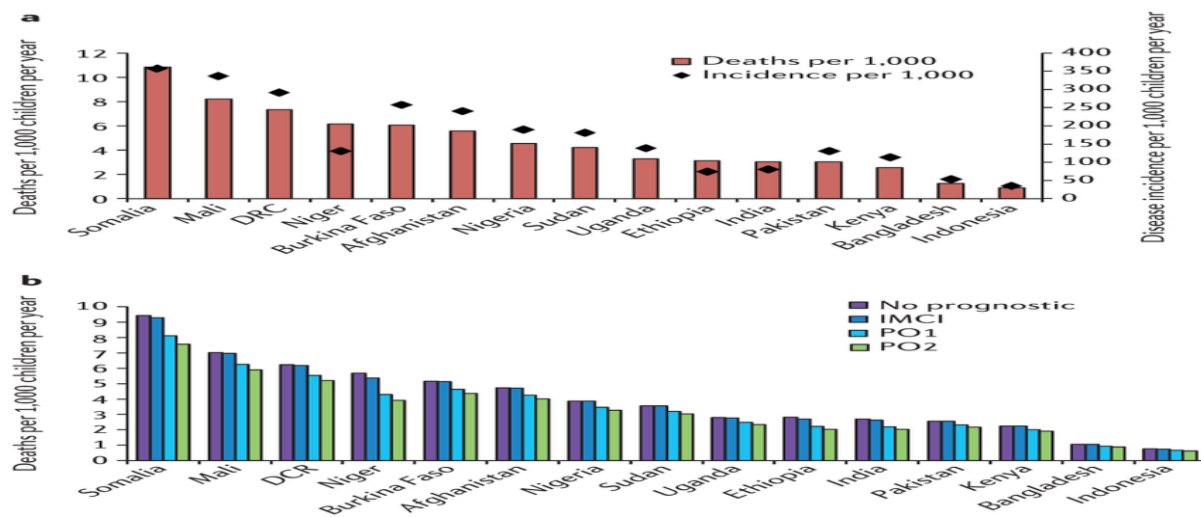
However, X-ray examination of the chest to detect pneumonia is a common subjective variation. Thus the default system for detecting pneumonia is required. In this study, we developed a **computer-assisted diagnostic system (CAD)** that uses a collection of study models of deep transmission with accurate classification of chest X-ray pictures.

In-depth learning is an important tool for practical wisdom, which plays an important role in it to solve many complex computer vision problems. In-depth learning models, in particular convolutional neural networks (CNNs), which are widely used to classify different images problems. However, such models only work well when given a large amount of data. With the problems of biomedical imaging problems, such a large number of labeled data is difficult to obtain because it requires specialist doctors to isolate each image which is an expensive and time-consuming task. Passing on learning is the task of overcoming this obstacle. In this process, to solve a problem involving a small database, the database-trained model is reused and network weights are determined in this regard. CNN models trained in large databases such as Image Net, which contains more than 14 million images, are often used for biomedical purposes.

Pneumonia is a respiratory illness caused by bacteria or viruses; it affects many people, especially in developing and developing countries. Pneumonia causes pleural leakage, a condition in which fluid leaks out fill the lungs, causing difficulty breathing. In this study, we developed a computer-assisted diagnostic system (CAD) that uses a collection of study models of deep transmission with accurate classification of chest X-ray pictures. Pneumonia is a serious respiratory disease that affects the lungs. It is ranked eighth in the list of top ten causes of death in the United States.

Early diagnosis of pneumonia is important to confirm therapeutic treatment and increasing survival rates. Pneumonia affects children and families everywhere but is most prevalent in South Asia and in sub-Saharan Africa. image classification of work. According to a study by Liu et al. in 2015, out of 5.9 million deaths of children under 5, more than 15.6% due to pneumonia; timely diagnosis and treatment can significantly reduce this mortality rate.

figure 2



However, the difference in chest X-ray images is low, making manual manipulation impossible. Computer-assisted diagnostics can improve efficiency and lead to timely treatment. The size, shape, and condition of pneumonia can vary greatly.

Its targeted content is unclear, which leads to greater detection difficulties, and increasing detection accuracy is a major research problem. Currently, screening algorithms include two-phase object detectors such as Faster R-CNN and single-stage detectors such as YOLO and SSD. The latter uses an additional section to complete the task of obtaining a high level target. They are faster than two-stage heifers but are less accurate. Medical tests have high accuracy requirements, and two-phase machines are advantageous in this regard.

Traditional deep networks reduce partial information when extracting features, and this affects their capacities for detection. The residual neural network structure of the low-complexity dilated bottleneck adopted in this paper avoids the computational lag associated with network depth and the problems involved in the large numbers of parameters associated with connectivity.

2.RELATED WORKS :

2.1 PNEUMONIA DETECTION WORKS

Pneumonia Detection is active in recent years, many experts have gone to great lengths to diagnose pneumonia. Abiyev and Maaitah used the convolutional neural network (CNN) to diagnose chest X-rays. Compared to BPNN and RNN, CNN gets high accuracy but long training time. Vijendran and Dubey combined multilayer over-the-counter (MLELM) and consecutive online learning (OSELM) machines to detect pneumonia in a chest X-ray image. Abiyev and Maaitah examined CNN's handwritten features and a set of classical features, including GIST and word bag in a database of more than 600 radiographs.

The above algorithms did well in detecting pneumonia, but the amount and size of the data involved was not large, and, in some cases, some experts used large amounts of data in a deep network to conduct research. Jaiswal et al predicted a possible pneumonia in the RSNA (Radiological Society of North America) database by Mask R-CNN, and the union-based mapping variance reached 21.8%. Gundel et al proposed using DenseNet to resolve the investigator on a chest X-ray database. Chakraborty et al. draw a convolutional neural network architecture consisting of a 17-layer network with multiple layers. Up to 95.62% AP on chest X-ray database. Wang et al. add visual acuity of multiple labeled labels with strong fragility and local disease framework to a deep neural convolutional network to solve problems on ChestX-ray8. The proposed methods are modified by network design, but not by spinal development. Only a spinal-directed spine is required.

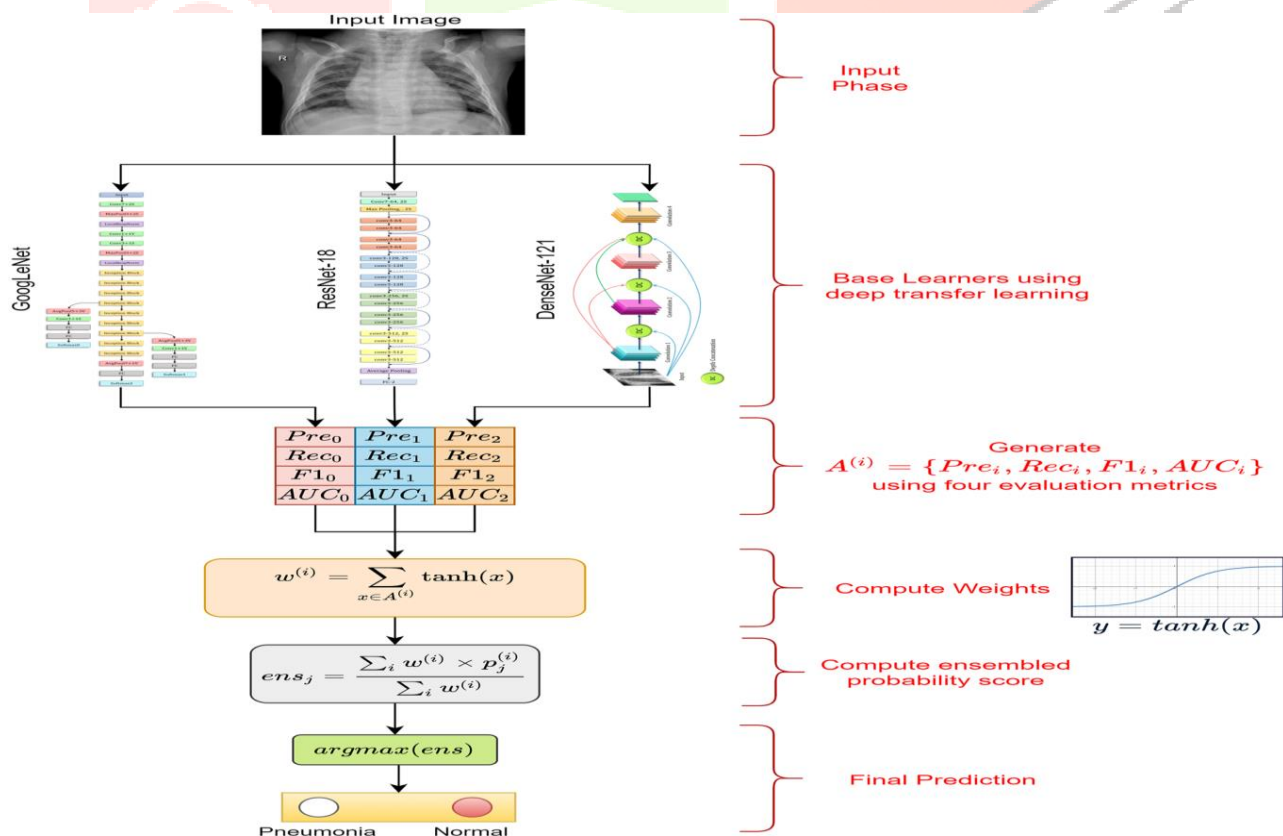
The COVID-19 pneumonia epidemic that broke out in late 2019 still threatens the survival of all humankind. At the same time, due to the rapid rate of new crown pneumonia infection, the rapid detection of crown pneumonia has placed significant demands on the global medical system. In-depth study helps in the diagnosis of new crowns.

Research on pneumonia has suggested a number of ways. In 2020, Wang and Wong proposed COVID-Net, a comprehensive neural network design designed to detect COVID-19 cases in chest X-ray images (CXR). Mangal Proposed CovidAID: COVID-19 AI Detector, which is a model based on a deep neural network to diagnose patients for appropriate diagnosis. Ozturk proposed a new model for the detection of COVID-19 using raw X-ray images of the chest; this model is used to provide an accurate diagnosis of binary categories (COVID vs. no findings) as well as multiple classifications (COVID vs. no findings vs. pneumonia). The first accuracy is 98.08%, and the latter is 87.02%. However, these studies performed only diagnostic tasks and eventually received X-ray diagnosis, and there was no way to obtain wound information.

2.2 PNEUMONIA DETECTION IN CHEST X-RAY IMAGES USING AN ESEMBLE OF DEEP LEARNING MODELS

figure 3 Representation of the proposed pneumonia screening framework. Previous = Accurate School, Rec = Memory School, F1 = F1-score, AUC = AUC school, and $A(i) = \{Pre_i, Rec_i, F1_i, AUC_i\}$; $w(i)$ the mass produced by the primary student to calculate the ensemble, points with the possibility of a j th sample by the i th editor, and end is a measure of the combined probability of a j th sample; and the function of argmax returns the highest value in the 1D system, that is, in this case it produces the predicted phase of the sample.

FIGURE 3



Diagnosis of pneumonia using chest X-rays has been an open problem for many years, the major limitation is the lack of publicly available data. Traditional machine learning methods have been extensively studied. Chandra et al [16] separated the lung regions from chest X-ray images and extracted eight mathematical features in these regions, which were used to isolate themselves. They used five traditional separators: a multi-layer perceptron (MLP), random forest, low sequential optimization (SMO), backward division, and setbacks. They tested their method in 412 images and obtained 95.39% accuracy using an MLP separator. Kuo et al [17] used 11 features to detect pneumonia in 185 patients with schizophrenia. They achieved a high level of accuracy, 94.5%, using the decision tree; some models are shorter with larger genes. Similarly, Yue et al. explained 6 factors used to detect pneumonia chest CT scan images of 52 patients; The highest number of AUC the amount they achieved was 97%. However, these methods cannot be generalized and tested in small databases. In contrast to machine learning algorithms, hand-created features need to be present, extracted and selected, classified or subdivided. Deep learning-based approaches make a distinction from end to end where appropriate and instructive features are present, automatically extracted from the input data and sorted.

CNN is preferred to image data separately because it automatically removes the fixed translation features by conversion of input image and filters. CNNs are flexible and efficient, better than the machine learning or common image processing techniques in image classification works and thus are widely used by researchers.

Sharma et al. [19] and Stephen et al. [20] have designed CNN's simple structures to distinguish X-ray images of lungs. Use data augmentation to compensate for data shortages. Sharma et al. obtained 90.68%, Stephen et al [20]. 93.73% accuracy and proposed a framework method based on enemy development in order to eliminate the reliance of the models on the source of the databases and produced solid speculation. Zhang et al. [23] used the machine learning-based approach in which they applied a subtle tree modification to images, followed by model classification. Then, extract the features using the local multikernel binary pattern and separate samples using traditional separators. They explored the path to a Small data comprising samples of COVID-19 and pneumonia also showed a positive effect with 97.01% accuracy. To solve the problem of data shortages in the activities of biomedical imaging activities, transfer learning, when using information obtained from large databases.

Table 2: Existing studies in pneumonia detection

METHOD	APPROACH	MERITS	DEMERITS
Chandra et al [17]	Lung x-rays segmentation using Image processing	95.39% accuracy achieved.	Limitation observed complex pattern recognition tasks
Kuo et al [17]	Traditional classifiers detailed examination in pneumonia detection	94.5% accuracy achieved.	Patient data are often private
Stephen et al [20]	7 layered CNN model implementation	93.78% accuracy achieved.	Computation cost high
Sharma et al [19]	Simple CNN model implementation	90.73% accuracy achieved	Computation cost high
Zhang et al [23]	Confidence aware module developed in x-ray scanning	97.01 accuracy achieved	Sensitivity obtained for dataset is low

3. PROPOSED METHODOLOGY:-

In this part we explain our proposed DeepConv-DilatedNet approach, which includes data processing, the design of our network, and the effective development effect of Soft-NMS in detail.

3.1 DATA PROCESSING

In 2018, the Radiological Society of North America (RSNA) [b20] released data on the diagnosis and localization of pneumonia on chest X-rays . The database is from National Health Centers (National Health Centers) Public chest X-ray images, with annotations by radiologist .Detailed data on the diagnosis of RSNA pneumonia can be found on the Kaggle website .The RSNA pneumonia data used in this trial contains 26684 cases data, of which only 6012 pneumonia images (approximately 22.03%), and the remaining 8851 normal images (approximately 31.19%) and 11821 images (accounting for 44.77%), is a rare or degenerative image of the lungs. In many in-depth readings, unintentional images do not help network training, so this insignificant piece of data is removed in the first step. Since a patient's chest pneumonia may have more than one area, there may be one to four areas. Therefore, in order to maintain the sample balance, we finally use 6012 images with annotations, of which 4/5 is selected as the training set and 1/5 as the test set.

3.2 PNEUMONIA DETECTION USING AN IMPROVED ALGORITHM BASED ON FASTER R-CNN METHOD

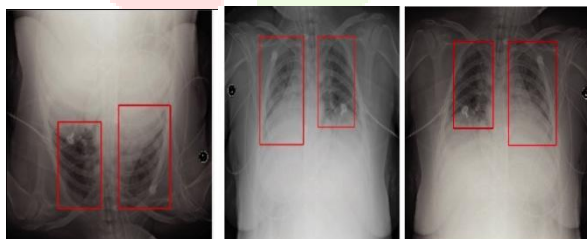
Table 3

Number of marking frames of lesions in the training set and test set.

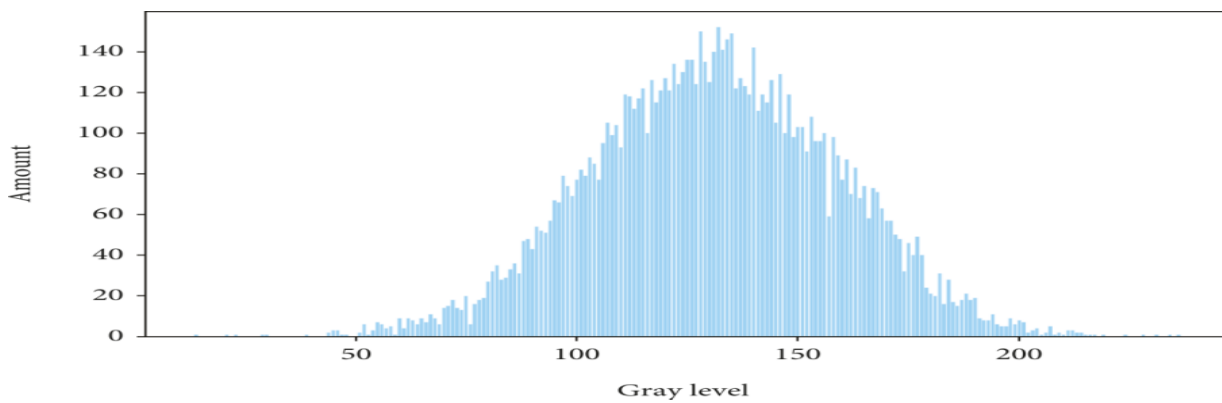
Dataset/lesion areas in each picture	1	2	3	4
Training set	2068	2628	104	10
Test set	545	629	25	3

figure 4

a) original image b) vertically flipped image c)horizontally flipped image



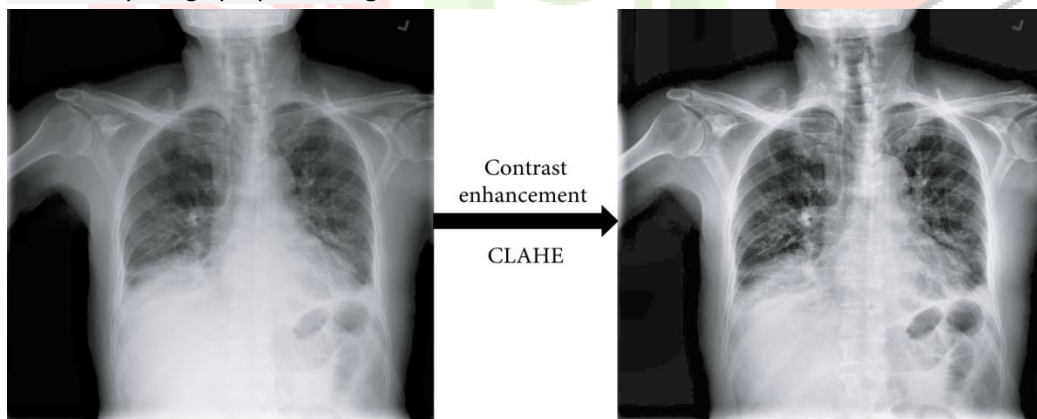
All 6012 images have pneumonia; among them, 3265 cases have two areas of pneumonia. Pneumonia is considered to be slightly more opaque than its surroundings. The analysis shows that the gray values are concentrated between 50 and 200, a very wide range. It is important to use the end-to-end method of deep learning.

figure 5

Due to the characteristics of low brightness and low contrast of the chest X-ray image, to better detect the target area of pneumonia and improve the detection accuracy, the chest X-ray image can be preprocessed. In X-ray images, normal lungs will not absorb X-rays, so it will appear black. The location of the pneumonia is a gray dashed shadow or a cloudy area.

To improve the recognition rate, contrast and brightness enhancement operations can be performed on X-ray images. In this article, the CLAHE algorithm is used to equalize the gray histogram to enhance contrast and brightness. To improve the effect of model training, it is necessary to enhance the training set. At first, we used the CLAHE algorithm to equalize the gray histogram of 6012 pictures with pneumonia given by RSNA, to enhance contrast and brightness. And then, the target area of pneumonia in every picture is used as a positive sample, and the remaining areas are used as a negative sample for training. Among them, 80% of samples, about 4810 pictures, are used as the training set, and 1202 pictures are used as the test set. The training set has less data, so it is increased by data enhancement. After horizontal flipping and vertical flipping, we finally get 14430 pictures as the training set. Figure 6 is shown below .

Chest X-ray image preprocessing



3.3 CONVOLUTION FUNCTION

The number of convolutional neural network statistics is shown in $N = ((W-F + 2P) \setminus S) + 1$ where N is the output size, W is the input size, F is the convolution kernel size, P is the padding value, and S is the step value. For example, we include an RGB image, our input image size is $227 * 227 * 3$, that is, it has three channels, and the size of each channel is $227 * 227$. We set the padding as 0 and the moving as 4, and then, the size of the convolution kernel is $3 * 3$. With the formula (1), we can calculate the output size as $N = (227-3 + 2 * 0) \setminus 4 + 1 = 57$. Generally, the larger the kernel of convolution, the larger the reception area, the more image information you can see, and the features you can get better. . Therefore, better features can be obtained by mixing convolution letters of different sizes. In this paper, $1 * 1$, $3 * 3$, and $7 * 7$ letters of convolution were used to obtain better feature maps.

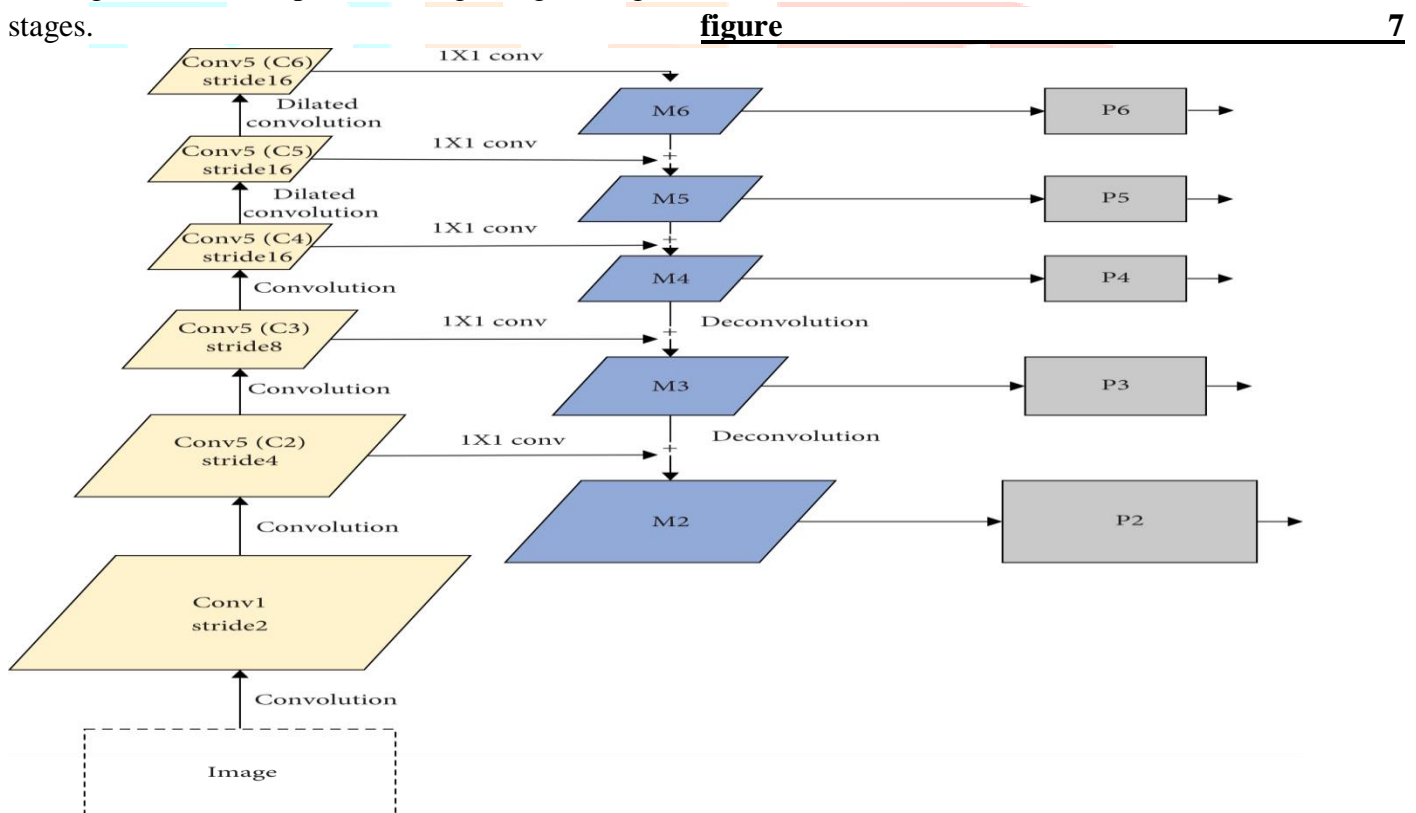
3.4 PNEUMONIA DETECTION USING DEEP LEARNING

The size of the anchor box will directly affect the performance of the model acquisition. This article analyzes the target data area of the pneumonia X-ray dataset. In the standard Faster R-CNN algorithm, the size of the anchor box depends on empiricism. It refers to the method of producing an anchor box in YOLOV 3 and uses the K-Means ++ algorithm to determine the appearance of the appropriate database size.

In this study, the K-Means ++ algorithm was used to analyze the target region of the training set, as well as the pneumonia box with three scales of [72, 73], [102, 120], and [140, 279]. . Therefore the scale of the anchor box was set to 0.5, 1.0, and 1.5.FPN combines low-level and high-end features to get sets of features that can display multidimensional information. Researchers use upsampling to locate a real-size feature map on high-resolution feature maps. Upsampling is an algorithm that restores an image to its original size. But deconvolution is another way to change the size of the feature maps.

The pneumonia-detection method used in this paper is based on the Faster R-CNN, in which the backbone uses the low-complexity dilated bottleneck residual neural network called DeepConv-DilatedNet. All the current state-of-the-art image classification networks are convolutional networks, such as ResNet and GoogLeNet, where the fully connected layer and upsample layers are replaced by convolution layers.

DeepConv-DilatedNet is a decomposing of the state-of-the-art image classification networks, such as ResNet and GoogLeNet. It removes the fully connected layer and only uses the convolutional layers to compute feature maps, speeding up the network. In DetNet, the first four phases of the backbone are consistent with the original ResNet50 phase, adding a stage, using convolutions and dilated convolutions in the fifth and sixth stages.



The RPN generates a large number of anchor boxes through the sliding window, with many overlapping parts between them. In this paper, the Soft-NMS literature is used to filter the overlapping anchor boxes, and it is hoped that this technique can be applied to computer screens as well.

3.5 SOFT -NMS

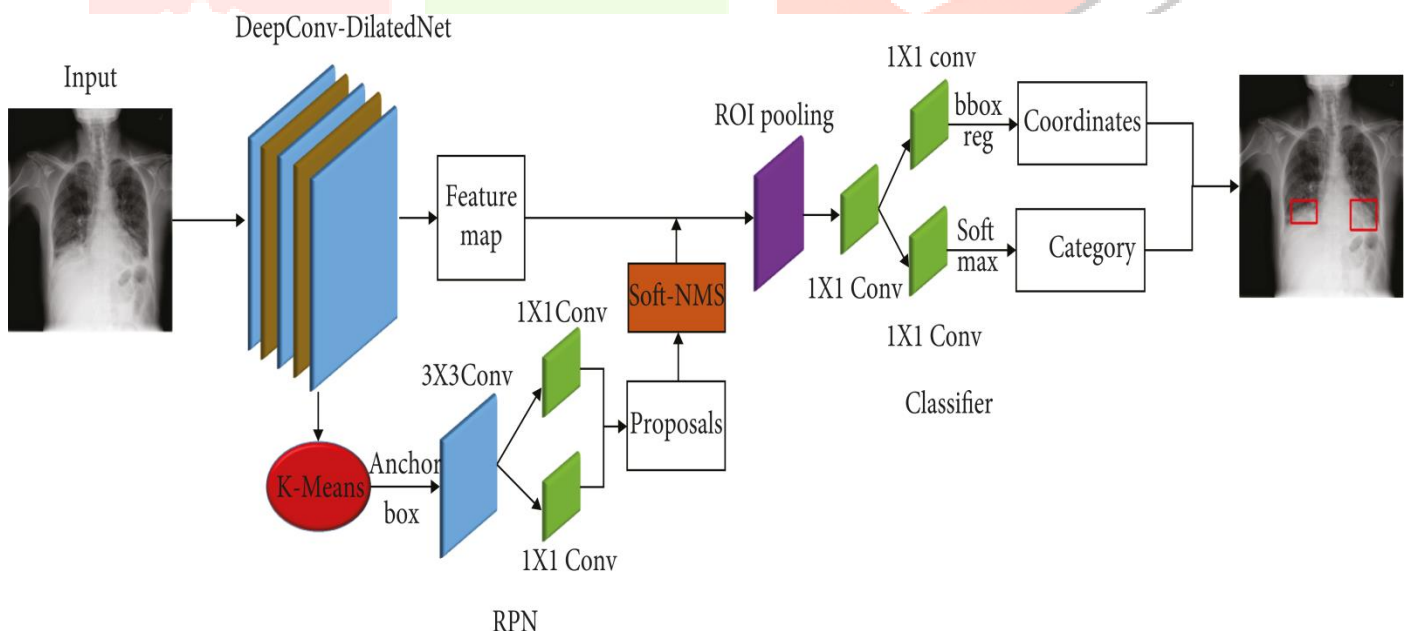
In the target acquisition process, whether using a slide window or RPN, multiple duplicate frames will be created.. The main idea is a repetitive finishing process, and a low score frame with a greater degree of overlap will be compressed by a high score frame. However, the traditional NMS approach uses strong decisions to determine which candidate structures are retained or compressed. So, something comes from a distance that is something else. That is, if two targeted frames are close together, a low-key frame will be ignored.

The spacing is too large and removable, causing object detection to fail and reducing the intermediate level detection algorithm. In this case, the acquisition algorithm should have two output acquisition frames, but the traditional non-standard value compression algorithm will be screened because the single-frame school is low and the IoU for two frames is larger than the set limit and Target detected. The test also shows that in one model, Soft-NMS can increase the effect of targeted acquisition from 39.8% to 40.9% .

4.METHOD ADOPTED

To find the target, first, we selected a two-segment R-CNN model with a suitable visual effect as a predictive framework, and then, consider the excellent DetNet59 effect as the backbone, so we used DetNet59 to extract the DetNet59 feature. intended. But because DetNet59 upsampling will lose features, we upgrade DetNet59 and convert upsampling into deconvolution to minimize feature loss, and at the same time, add hole convolution to extend the reception field. Apart from that, the first ancho box had a huge impact on model training and prediction, so we borrowed the YOLO method and used the K-Means ++ algorithm to obtain the first ancho box of targeted pneumonia databases to improve the detection effect. Finally, because Soft-NMS improves network performance for targeted acquisition, we have joined Soft-NMS. Finally, we get DeepConv-DilatedNet + Faster R-CNN which is shown as Figure 8 below.

figure 9

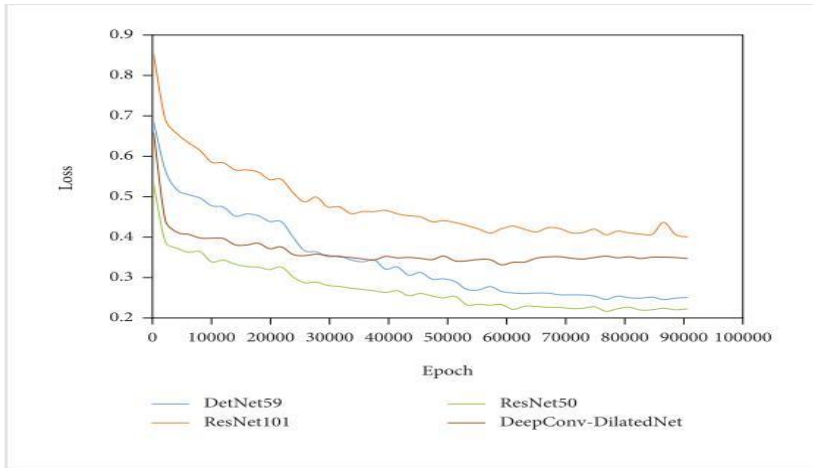


4.1 PROCEDURE:

Initially, we did not filter the original RSNA data set, so the database is full of many useless images, and the result of a trained model using such a database is very bad. After data analysis, the data without data development was used to train the model, but because the data value was too small, the model could not fully integrate and the prediction result was not good. Finally, we re-evaluated the database and obtained the training results of the current model.

The test was performed on the NVIDIA GeForce GTX 1080 configuration. With Cuda acceleration, 2 images are processed per set, reading rate is set at 0.001, and period 10 is reduced tenfold using the SGD optimization algorithm. The retreat of the RPN binding box uses the Smooth L1 loss function, and the split uses the binary loss function (CE). To ensure performance, we have trained and evaluated four different lines. The repetition value exceeds 90000, and the model appears stable. We plan to split and reverse the total loss curve for each model shown in Figure 7. The DeepConv-DilatedNet loss rate varies by a range

0.4 to 0.5. The correct result is correct. The loss rate for DetNet59 and ResNet50 is steady around 0.24.



The smooth drop of VGG16 eventually stabilized at 0.25; the ResNet101 fluctuations are slightly stable at about 0.45, with a negative impact.

4.2 RESULT EVALUATION

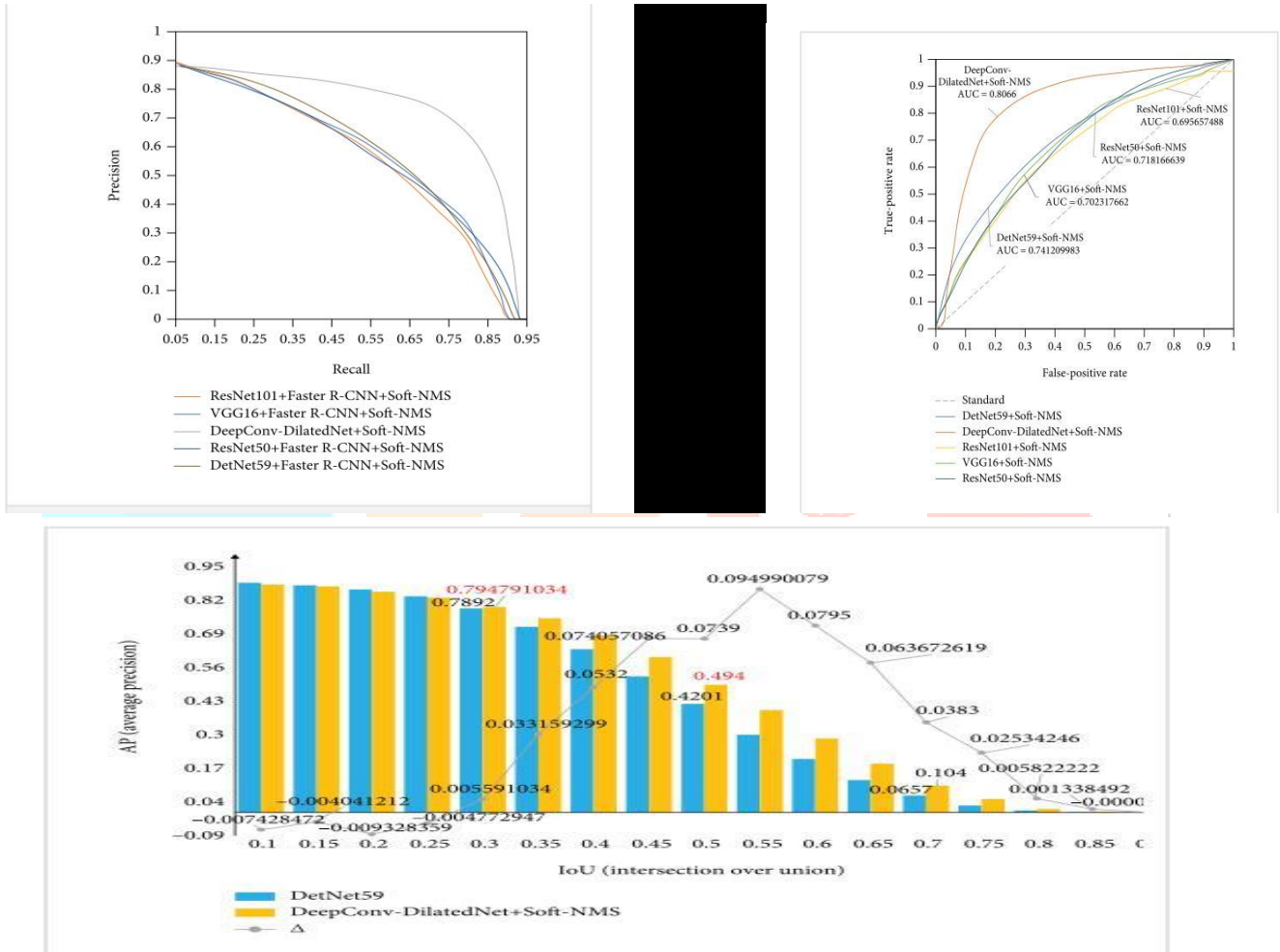
A value based on the IoU-threshold-based Average Precision (AP) is used to indicate acquisition results. IoU is the level of discrepancy between the predictive box and the basic truth box. As can be seen in Formula (1), the AP refers to the central points of each image. $AP(s,i) = \sum i s_i / N$ -----(1)

Network	mAP@0.4	mAP@0.5	mAP@0.6	mAP@0.7	mAP
DetNet59+Soft-NMS	0.6617	0.4751	0.2638	0.0879	0.3721
DeepConv-DilatedNet+Soft-NMS	0.6849	0.4940	0.2863	0.1040	0.3923
ResNet50+Soft-NMS	0.6234	0.4268	0.2495	0.0842	0.3460
ResNet101+Soft-NMS	0.5790	0.3992	0.2077	0.0630	0.3122
VGG16+Soft-NMS	0.5925	0.4179	0.2462	0.0940	0.3377

Assessment results for different IoU thresholds. $IoU(Pred, GT) = \frac{Pred \cap GT}{Pred \cup GT}$ -----(2)

In Formula (2), Pred is a prediction box, GT refers to the basic truth box, in which it presents the points of each test image and specifies the number of all test images. Suppose an IoU threshold is equal to a certain limit where the IoU limit of the predicted image is above or equal to the limit; otherwise, it is 0. $Map(p, t) = \sum i p_i / n$ -----(3).

The Mean Average Precision (mAP) is used to describe the acquisition accuracy of each model, such as Formula (1), the IoU limit and the AP value at the border by refers to the number of classes, so in this paper the ROC is a tool for measuring inequality in classification [24], and its abscissa is the false positive (FPR) level as indicated in Formula (4), and the ordinate is the positive-positive value (TPR) as indicated in the Formula 5. ROC curves are often used to evaluate the pros and cons of a binary category. When the ROC curve approaches the upper left corner, the true value of the detector obtained is much higher compared to its false level, indicating that the separator is performing well. $PR = FP \setminus (FP + TN)$ $TPR = TP \setminus (TP + FN)$



Assessment results for(soft NMS4, 5)

	AP@0.4	AP@0.5	AP@0.6	AP@0.7	mAP
DetNet59	0.6317	0.4201	0.2068	0.0657	0.3311
ResNet50	0.6066	0.3791	0.1863	0.0513	0.3058
ResNet101	0.5539	0.3508	0.1540	0.0406	0.2748
VGG16	0.5506	0.3559	0.1881	0.0660	0.4210
DeepConv-DilatedNet	0.6419	0.4570	0.2732	0.0746	0.3617

To fully validate the test results, we calculated the value of the AUC (lower area curve) [25]. AUC is a number of opportunities. If positive and negative examples are randomly selected, chances are that the current classification algorithm measures positive conditions before negative based on the calculated result of the AUC value. Therefore, the greater the number of AUCs, the greater the likelihood that a segmentation algorithm will weigh positive examples before negative ones, making better segmentation better.

4.3. Test Result

In the experiments, comparisons of four different AP spinal indicators with different IoU limits were performed, and mAP () values were calculated as shown in Table 2. First, comparing the model acquisition effect using DetNet59 as the backbone and method used. Vgg16, ResNet50, and ResNet101 as the backbone, respectively, found that the result of using DetNet59 as a spinal model is much better than others. Then, we compare our advanced network with DetNet59; the result shows that our method is better than DetNet59. As can be seen in Table 2, Faster R-CNN with DeepConv-DilatedNet as the core is higher than DetNet59, ResNet50, ResNet 101, and VGG16 networks in each AP with different layers. In the RSNA database, DeepConv-DilatedNet reached 0.4
Table 4: Comparison of results for different networks. Network MS Mask R-CNN 0.2181 DeepConv-DilatedNet+Soft-NMS 0.35087 Assessment results for soft NMS.

To understand the usefulness of our model, we test our method on the ChestX-ray14 dataset. The ChestX-ray14 dataset contains 30805 patients and 112,120 chest X-ray images. The size of each image is with 8 bits grayscale values. The corresponding report includes 14 pathology classes. There are 120 pneumonia images with bounding box annotations in ChestX-ray14. We choose all of them to check our model. we can see the detection effect of DeepConv-DilatedNet is still better than other models.

CONCLUSION:- Early detection of pneumonia is important in determining the appropriate treatment for the disease and preventing it from endangering the patient's health. Chest radiographs are the most common a widely used tool for diagnosing pneumonia; however, they are subject to category variation and the diagnosis is based on the doctors' expertise in diagnosing pneumonia. To help physicians, an automated CAD system was developed in this study, which uses in-depth reading-based classification to divide chest X-ray images into two classes "Pneumonia" and "Common." An integration framework was developed that looked at the decision points obtained from the three CNN models, GoogLeNet, ResNet-18, and DenseNet-121, so that create a limited integrated reading. In this paper, a complex low-level neural network with an expanded bottle structure, called DeepConv-DilatedNet, is used as the backbone of a two-phase detector using Faster R-CNN. Due to the variability of the pneumonia target, the image has also been enhanced with the CLAHE algorithm to make the target area more prominent. At RPN, we use the Soft-NMS algorithm to filter the anchor box and verify its quality. In order to speed up the algorithm integration and improve the accuracy of the target location, we also used the K-Means ++ algorithm in YOLOV3 to determine the initial size of the anchor box. We invest de convolutions in FPN on scale variations and thus facilitate recognition from features listed on a single input scale. Finally, we got the result of this approach. Combining different sets of work done on each network, the ability of the pneumonia algorithm to accurately detect RSNA databases is enhanced. To ensure the validity of the model, we also compared it in detail with the traditional networks of DetNet59, ResNet50, ResNet101, and VGG16 and compared it with other high quality results; Our algorithm does a good job at this task.

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