

DETECTION OF COVID-19

D. Chaithanya
Assistant Professor
Department of Information Technology
Vignan Institute of Technology and Science
Hyderabad, India
dchaitu94@gmail.com

Goli Deepa
UG Scholar
Department of Information Technology
Vignan Institute of Technology and Science
Hyderabad, India
golideepareddy1122@gmail.com

Kondapalli Bindu Pavani
UG Scholar
Department of Information Technology
Vignan Institute of Technology and Science
Hyderabad, India
bindukondapalli28@gmail.com

Gunture Manjusha
UG Scholar
Department of Information Technology
Vignan Institute of Technology and Science
Hyderabad, India
gunturemanjusha2001@gmail.com

Vangari Vishal Kumar
UG Scholar
Department of Information Technology
Vignan Institute of Technology and Science
Hyderabad, India
vvishalk9505@gmail.com

ABSTRACT

COVID-19 global pandemic affects health care and lifestyle worldwide, and its early detection is critical to control cases' spreading and mortality. The actual leader diagnosis test is the Reverse transcription Polymerase chain reaction (RT-PCR), result times and cost of these tests are high, so other fast and accessible diagnostic tools are needed. Inspired by recent research that correlates the presence of COVID-19 to findings in Chest X-ray images, this paper's approach uses existing deep learning models (VGG19 and U-Net) to process these images and classify them as positive or negative for COVID-19. The proposed system involves a preprocessing stage with lung segmentation, removing the surroundings which does not offer relevant information for the task and may produce biased results; after this initial stage comes the classification model trained under the transfer learning scheme; and finally, results analysis and interpretation via heat maps visualization. The best models achieved a detection accuracy of COVID-19 around 97%.

I. INTRODUCTION

Coronavirus infection is caused by Severe Acute Respiratory Syndrome (SARS) and the Middle East Respiratory Syndrome (MERS). The illness induced by SARS-CoV-2 is caused by a new coronavirus, COVID-19 (Zhang, 2020). The first COVID-19 cases were reported in December 2019 in Wuhan, Hubei Province, China (Xu et al., 2020). On March 11, 2021, the World Health Organisation (WHO) designated COVID-19 a pandemic (Ducharme, 2020), and there have been 188,404,506 documented cases worldwide, resulting in 4,059,220 fatalities (Worldometer, 2020).

MOTIVATION

These disorders produce respiratory issues that are treatable without the need of specialised medication or equipment. Nonetheless, underlying medical conditions such as cancer, cardiovascular disease, diabetes, and respiratory illness might exacerbate this illness (World Health Organisation, 2020).

Transcription in reverse The major approaches for COVID-19 detection are currently a polymerase chain reaction (RT-PCR) and gene sequencing for respiratory or blood samples (Wang et al., 2020).

2. LITERATURE SURVEY

This study introduces a novel technique that makes use of current Deep Learning models. It focuses on improving the preprocessing step in order to generate accurate and reliable COVID19 classification results from Chest X-ray images. The preprocessing step includes a network to filter the images based on the projection (lateral or frontal), some common operations such as normalisation, standardisation, and resizing to reduce data variability, which may affect the performance of the classification models, and a segmentation model (U-Net) to extract the lung region that contains the relevant information while discarding the information from the surroundings that can produce misleading results (de Inform). Following the preprocessing step, the classification model (VGG16-19) is used, which employs the transfer learning technique, which makes use of pre-trained weights from a much larger dataset.

3.PROPOSED SYSTEM

In a proposed system,the work can be automatically done by Automation tool without any human intervention.

The proposed system involves a preprocessing stage with lung segmentation, removing the surroundings which does not offer relevant information for the task and may produce biased results; after this initial stage comes the classification model trained under the transfer learning scheme; and finally, results analysis and interpretation via heat maps visualization. The best models achieved a detection accuracy of COVID-19 around 97%.

4. SYSTEM ARCHITECTURE

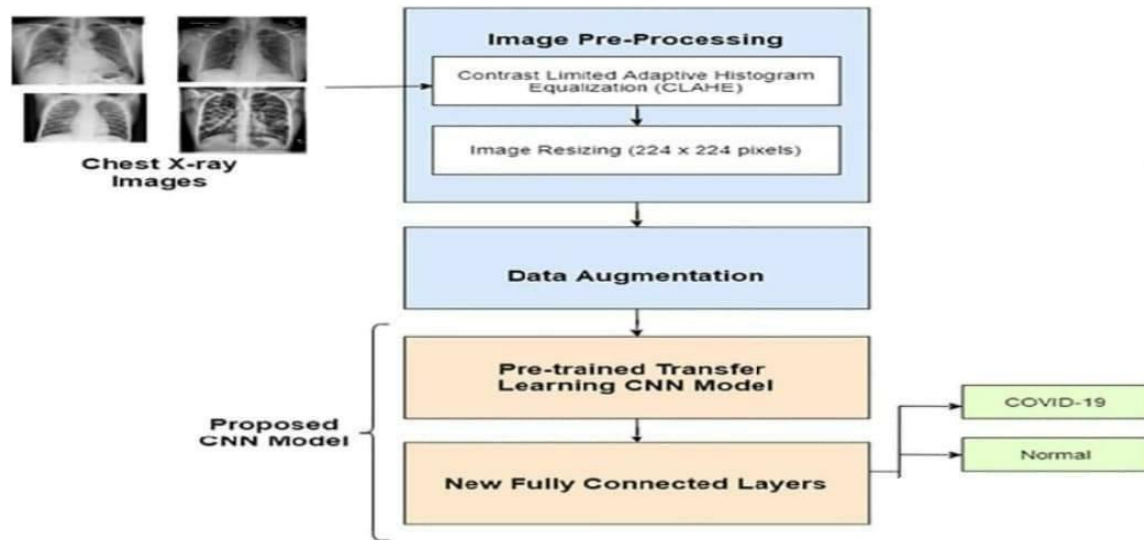


Fig 1. System Architecture for covid-19 detection

Data Collection

Data is gathered from several sources at this level. Data collection is the process of obtaining and analysing information from several sources. The type of data gathered is determined by the targeted project. Data may be gathered from a variety of sources.

Sources such as files, databases, etc. The accuracy of the intended system is directly affected by the quality and amount of data obtained. Predictive models are only as good as the data on which they are based, hence excellent data collecting practises are essential for constructing high-performing models.

Data Preprocessing

Data preparation is done at this step to sanitise the raw data. Data acquired in the actual world is cleaned up. Missing values, inconsistent values, duplicate instances, and so on may be present in raw data. As a result, raw data cannot be utilised directly to develop a model.

There are several approaches for cleaning the dataset.

- Ignoring the values that are missing
- Removing missing-value instances from the dataset. Estimating missing-value instances using mean, median, or mode.
- Duplicate instances are removed from the dataset.

Choosing learning

Algorithm The best performing learning algorithm is studied at this stage. It depends on the type of problem and the data we have. Classification algorithms are employed when the problem is to classify and the data has been labelled. Regression algorithms are employed when the goal is to execute a regression job on labelled data. Clustering methods are employed if the objective is to form clusters and the data is unlabeled.

Training model

The model is being trained at this stage to increase its capabilities. The dataset is split into two parts: training and testing. The training/testing balance is around 80/20 or 70/30. It is also determined by the amount of the dataset. The training dataset is utilised to train.

A testing dataset is used for testing. The learning algorithm receives the training dataset. The Learning Algorithm builds the model by discovering a mapping between the input and the output.

Evaluating model

The model is assessed at this level to see how good it is. The retained-aside testing dataset is used to assess the model. It enables the model to be tested on data that has never been used for training before. To assess performance, metrics like as accuracy, precision, recall, and so on are utilised. If the model fails to perform adequately, it is rebuilt using new hyper parameters. The precision may be enhanced even further by adjusting the hyper settings.

5. WORKING

Install all the required software requirements

Open Anaconda Prompt and Activate the project by running the following commands:

```
cd path(paste the copied path)
```

```
conda activate project
```

```
Spyder
```

After running these commands, it will redirect to Spyder containing the code required to execute our project.

By selecting and loading the path of any image in the testcases of the code, By clicking enter the result can be predicted whether it is covid positive or negative.

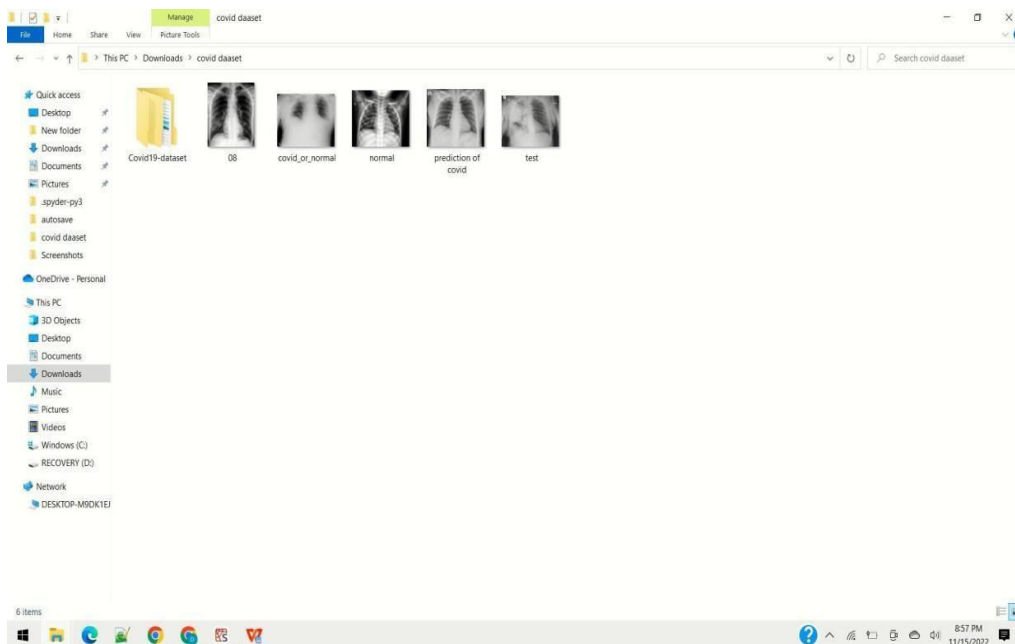


Fig.2: The folder consists of Types of datasets

Now run all the cells in the Spyder.

All the cells will be executed one after the other and produces the required output.

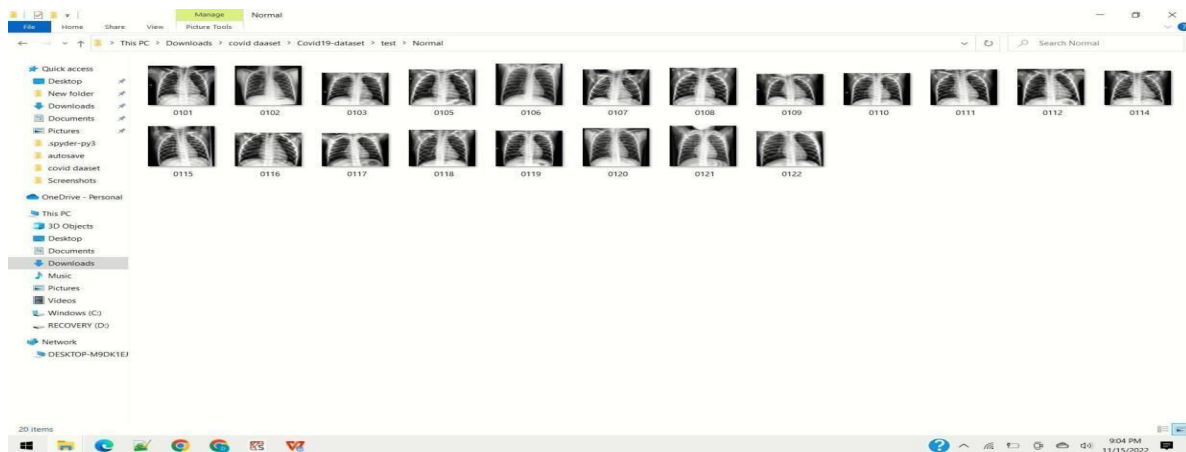


Fig 3. All X-ray images in dataset

A screenshot of the Spyder Python IDE console. The console shows the execution of a Python script for X-ray image classification. The script imports numpy, tensorflow, and keras, then loads an X-ray image, preprocesses it, and uses a trained model to predict the class. The output shows the prediction for two different images: 'COVID NEGATIVE' and 'COVID POSITIVE'. The console also shows the execution time for each step: 78ms/step and 109ms/step. The taskbar at the bottom shows the system tray with the date and time: 5:46 PM, 11/15/2022.

Fig 4. Prediction of results of X-ray

6. CONCLUSION

This technique demonstrates how old models may be useful for many tasks, even when the updated U-Net modelling do not perform better. It is also demonstrated how visual noise might cause bias in the computational models. The photos without segmentation are superior for identifying COVID illness according to most measures. Further investigation reveals that, even if the metrics are improved, these models are predicated on observable disorders across the lungs as evident proof of COVID, implying that true correct models must centre on lung sections for classification. In this scenario, segmentation is required to provide trustworthy findings by eliminating bias. Transfer learning was critical to the outcomes provided.

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