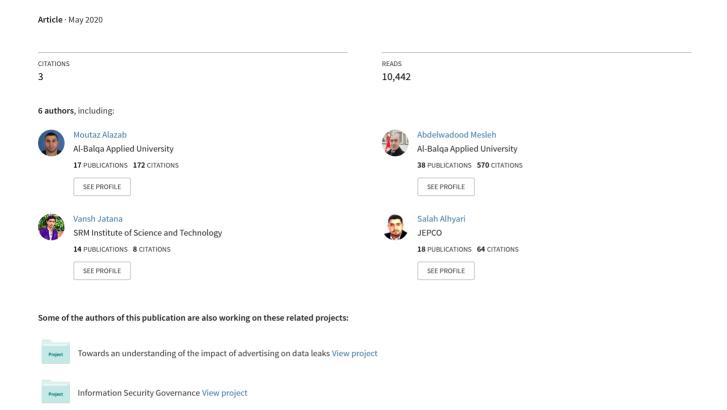
## COVID-19 Prediction and Detection Using Deep Learning



Received: 27 April 2020; Accepted: 28 May, 2020; Published: 29 May, 2020

# COVID-19 Prediction and Detection Using Deep Learning

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Abstract: Currently, the detection of coronavirus disease 2019 (COVID-19) is one of the main challenges in the world, given the rapid spread of the disease. Recent statistics indicate that the number of people diagnosed with COVID-19 is increasing exponentially, with more than 1.6 million confirmed cases; the disease is spreading to many countries across the world. In this study, we analyse the incidence of COVID-19 distribution across the world. We present an artificial-intelligence technique based on a deep convolutional neural network (CNN) to detect COVID-19 patients using real-world datasets. Our system examines chest X-ray images to identify such patients. Our findings indicate that such an analysis is valuable in COVID-19 diagnosis as X-rays are conveniently available quickly and at low costs. Empirical findings obtained from 1000 X-ray images of real patients confirmed that our proposed system is useful in detecting COVID-19 and achieves an F-measure range of 95-99%. Additionally, three forecasting methods—the prophet algorithm (PA), autoregressive integrated moving average (ARIMA) model, and long short-term memory neural network (LSTM)-were adopted to predict the numbers of COVID-19 confirmations, recoveries, and deaths over the next 7 days. The prediction results exhibit promising performance and offer an average accuracy of 94.80% and 88.43% in Australia and Jordan, respectively. Our proposed system can significantly help identify the most infected cities, and it has revealed that coastal areas are heavily impacted by the COVID-19 spread as the number of cases is significantly higher in those areas than in non-coastal

*Keywords*: Artificial Intelligence, X-ray, Convolutional Neural Network, Machine Learning, COVID-19.

#### I. Introduction

The coronavirus disease (COVID-19) is a global pandemic that was discovered by a Chinese physician in Wuhan, the capital city of Hubei province in mainland China, in December 2019 [1]. Currently, there is no approved human

vaccine for combating it. COVID-19 propagation is faster when people are in close proximity. Thus, travel restrictions control the spread of the disease, and frequent hand washing is always recommended to prevent potential viral infections. Meanwhile, fever and cough are the most common infection symptoms. Other symptoms may occur, including chest discomfort, sputum development, and a sore throat. COVID-19 may progress to viral pneumonia which has a 5.8% mortality risk. The death rate of COVID-19 is equivalent to 5% of the death rate of the 1918 Spanish flu pandemic.

The total number of people infected with COVID-19 worldwide is 5,790,103 as of May 27, 2020 whereas the numbers of reported deaths and recoveries are 357,432 and 2,497,618 respectively. Most of the cases were recorded in the USA, Spain, Italy, France, Germany, mainland China, UK, and Iran [2]. Saudi Arabia, with 78,541 cases, has the highest number of reported cases among all the Arab countries. Meanwhile, the number of reported cases in Jordan is 720, whereas the numbers of deaths and recoveries are 9 and 586 respectively. The number of reported cases in Australia is 7150, whereas the numbers of deaths and recoveries are 103 and 6579, respectively. Since February 2020, information technology services, such as mobile apps, have been used to curb the potential risk of infection in mainland China. The mobile apps suggest users to self-quarantine and alert the concerned health authorities when someone infected by the virus. They also monitor infected people, and the last persons that they had contact with [3].

Since it was first reported, the disease has spread exponentially across the world and has become an international concern. A research conducted by Jiang et al. [4] revealed that the death rate of COVID-19 is 4.5% across the world. The death rate of patients in the age range of 70–79 years is 8.0%, whereas that of patients above 80 years is

14.8%. The authors also confirmed that patients above the age of 50 years with chronic illnesses are at the highest risk and should therefore take special precautions. One of the main threats of COVID-19 is its rapid propagation, with an estimated 1.5–3.5 people getting infected by the disease upon contact with an infected person [5]. This implies that if 10 people are COVID-19 positive, they are more likely to infect 15–35 other people. Therefore, COVID-19 can infect a very large number of people in a few days unless intervention measures are implemented.

The standard diagnostic technique is the reverse transcription-polymerase chain reaction (RT-PCR) method [1], a laboratory procedure that interacts with other ribonucleic (RNA) and deoxyribonucleic acids (DNA) to determine the volume of specific ribonucleic acids using fluorescence. RT-PCR tests are performed on clinical research samples of nasal secretions. The samples are collected by inserting a swab into the nostril and gently moving it into the nasopharynx to collect secretions. Although RT-PCR can identify the severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) strain that causes COVID-19, in some cases, it produced negative test results even though the patients showed progression on follow-up chest computed tomography (CT) scans [6]. In fact, several studies [6-9] have recommend the use of CT scans and X-rays rather than RT-PCR owing to its limited availability in some countries.

The detection of COVID-19 symptoms in the lower parts of the lungs has a higher accuracy when using CT scans or X-rays than that when using RT-PCR [7]. In certain cases, CT scans and X-ray tests can be substituted with RT-PCR tests. However, they cannot exclusively address the problem owing to the relatively limited number of radiologists, compared to new residents, and the high volume of re-examinations of infected people who wish to know the progression of their illness. To overcome the challenges of CT scans and X-rays and to assist radiologists, we need to improve the speed of the procedure. This can be achieved by designing advanced diagnostic systems that utilise artificial intelligence (AI) tools. The aim is to reduce the time and effort required to perform CT scans and X-rays of COVID-19-positive patients and evaluate the rate of disease development [7-9].

Radiological imaging is considered an important screening method for COVID-19 diagnosis [10]. Ai et al. [6] demonstrated the consistency of the radiological history of COVID-19-related pneumonia with the clinical nature of the disease. When examined by CT scans, almost all COVID-19 patients have exhibited similar features including ground-glass opacities in the early stages and pulmonary consolidation in the latter stages. In fact, the morphology and peripheral lung distribution can be rounded [6]. AI can be used to initially evaluate a COVID-19 patient as an alternative solution to traditional approaches that are time-consuming and labour-intensive. In this paper, we advocate the use of AI to forecast COVID-19 cases and diagnose COVID-19 patients via chest X-ray images.

#### A. Contributions of This Study

The following are the core contributions of this study:

• We propose an automated intelligent system for distinguishing COVID-19 patients from non-patients on the basis of chest X-ray images. Our system instantly reads the structure of a chest X-ray image, leverages hidden patterns to identify COVID-19 patients, and reduces the need for manual pre-processing steps.

- Empirical findings obtained from 1000 chest X-ray images of patients confirmed that our proposed system can detect COVID-19 patients with an accuracy of 95–99%.
- We provide an intelligent prediction system for predicting the number of patients confirmed to have contracted the disease, recovered from the disease, and died from the disease over the next 7 days using three forecasting methods. Our proposed system has been trained and tested on datasets generated from real-world cases and has predicted the numbers of COVID-19 confirmations, recoveries, and deaths in Australia and Jordan with an average accuracy of 94.80% and 88.43%, respectively.
- We highlight the most affected areas and show that coastal areas are heavily impacted by COVID-19 infection and spread as the number of cases in those areas is significantly higher than that in other non-coastal areas.

The rest of this paper is organised as follows. Section 2 presents the related works on recent COVID-19 detection and prediction methods for chest X-ray images. Section 3 presents the detailed system design, dataset description, and performance-evaluation metrics. Sections 4 and 5 present the results and discussions, respectively. Section 6 concludes the paper and provides an outlook to future research.

#### **II. Related Works**

The analysis and detection of COVID-19 have been extensively investigated in the last few months. The first part of this section addresses issues related to COVID-19 detection based on deep-learning approaches using CT scans and chest X-ray images. The second part reviews the related literatures to assess future estimates of the number of COVID-19 confirmations, recoveries, and deaths.

COVID-19 has now become a global pandemic owing to its rapid spread. It is very challenging to detect exposed persons because they do not show disease symptoms immediately. Thus, it is necessary to find a method of estimating the number of potentially infected persons on a regular basis to adopt the appropriate measures. AI can be used to examine a person for COVID-19 as an alternative to traditional time-consuming and expensive methods. Although there are several studies on COVID-19, this study focused on the use of AI in forecasting COVID-19 cases and diagnosing patients for COVID-19 infection through chest X-ray images.

Several research areas have implemented AI (e.g. disease diagnoses in healthcare) [11-13]. One of the main advantages of AI is that it can be implemented in a trained model to classify unseen images. In this study, AI was implemented to detect whether a patient is positive for COVID-19 using their chest X-ray image.

AI can also be used for forecasting (e.g., how the population will increase over the next 5 years) through existing evidence. Thus, predicting possibilities in the immediate future can help authorities to adopt the necessary measures [14]. Wynants et al. [15] focused on two main concepts. The first concept involved studies related to the diagnosis of COVID-19, and the second involved studies related to the prediction of the number of people who will be infected in the coming days. The study analysis maintained that most of the existing models are poor and biased. The authors suggested that research-based COVID-19 data should be publicly available to encourage the adoption of more specifically designed detection and prediction models.

#### **COVID-19 Diagnosis Using Deep Learning**

The use of machine learning (ML) has been rapidly increasing in various fields including malware detection [16-19], mobile malware detection [20-24], medicine [25-27] and information retrieval [27-31]. In 2012, a modern ML system called deep learning was introduced, which is based on a convolutional neural network (CNN). It won the ImageNet classification competition, the world's best-known computer-vision Deep-learning algorithms competition [32]. computational models composed of multiple processing layers to learn data representation through several abstraction layers. They train a computer model to perform classification tasks directly from pictures, texts, or sounds. According to LeCun et al. [33], deep-learning models feature high accuracies and can improve human output in certain instances.

#### X-Ray Diagnosis Using Deep Learning

X-ray machines use light or radio waves as radiation to examine the affected parts of the body because of cancers, lung diseases, bone dislocations, and injuries. Meanwhile, CT scans are used as sophisticated X-ray machines to examine the soft structures of active body parts for better views of the actual soft tissues and organs [34]. The advantages of using X-rays over CT scans are that X-rays are quicker, safer, simpler, and less harmful than CT scans [7, 34].

Narin et al. [7] proposed a CNN-based model to detect COVID-19 patients using 100 chest X-ray images, half of which belong to COVID-19 patients and the other half belong to healthy people. They evaluated three CNN models—ResNet-50, Inception-v3, and Inception-ResNet-v2—using five-fold cross-validation and reported that ResNet-50 had the best detection accuracy (98%).

In a similar study conducted by Sethy and Behera [35], the authors extracted features from chest X-ray images using a deep-learning algorithm and classified the images as either infected or healthy using a support vector machine (SVM). The authors employed 11 deep-learning models: AlexNet, VGG16, VGG19, GoogLeNet, ResNet-18, ResNet-50, ResNet-101, Inception-v3, Inception-ResNet-v2, DenseNet-201, and XceptionNet. They collected two datasets—the first containing chest X-ray images of 25 infected patients and 25 non-infected patients and the other containing chest X-ray images of 133 infected patients (e.g. MERS, SARS, and ARDS patients) and 133 non-infected patients. They performed separate feature extractions on each dataset using various models and achieved a 95.38% accuracy with ResNet-50 and SVM.

Furthermore, Hemdan et al. [36] proposed a framework, called COVIDX-Net, that can assist radiologists in diagnosing COVID-19 patients using X-ray. They evaluated their framework using a dataset of 50 X-ray images divided into two classes: 25 COVID-19-positive images and 25 COVID-19-negative images. The images used were resized to 224×224 pixels. The COVIDX-Net framework employs seven deep learning models: MobileNet, ResNet-v2, Inception-ResNet-v2, Xception, Inception-v3, DenseNet, and modified VGG19. Their evaluation results indicate that the VGG19 and DenseNet models delivered comparable performances with an F-score of 91% for COVID-19 cases.

In addition, Hassanien et al. [37] proposed a classification system that uses multi-level thresholding and an SVM to detect COVID-19 in lung X-ray images. Their system was tested on 40 contrast-enhanced lung X-ray images (15 healthy and 25 COVID-19-infected regions) with a resolution of

 $512 \times 512$  pixels. Their classification system achieved a sensitivity of 95.76%, a specificity of 99.7% and an accuracy of 97.48%.

#### 1) CT Scan Diagnosis Using Deep Learning

The CT scan was developed by Godfrey Hounsfield and Allan Cormack in 1972. It utilises an advanced X-ray technology to carefully diagnose delicate internal organs [34]. CT scanning is quick, painless, non-invasive, and precise and can produce three-dimensional (3D) images [34]. CT scans of internal organs, muscles, soft tissues, and blood vessels offer greater clarity than standard X-rays, especially for soft tissues and blood vessels. The main disadvantage of the CT scan is that it is expensive, compared to X-rays [34].

The sensitivity and specificity of RT-PCR for COVID-19 detection have been criticised in several studies [4, 38, 39]. Although RT-PCR is the standard method for this purpose, it generates a relatively large number of false negatives owing to several reasons, including methodological drawbacks, disease stages, and methods of obtaining the specimens, which delay disease diagnosis and control. Therefore, RT-PCR tests are not sufficient for assessing the disease status. Recent results have revealed that nucleic acid testing is not reliable and can only achieve an accuracy of 30–50% [38].

Jiang et al. [4] compared RT-PCR to CT scans and examined 51 patients (29 men and 22 women) with a history of travel to or residency in endemic areas and with severe respiratory and fever symptoms due to unknown causes. The authors obtained a sensitivity of 98% in a non-contrast chest CT scan for the detection of COVID-19, compared to the initial RT-PCR sensitivity of 71%. Owing to the shortage of RT-PCR kits and the growing number of COVID-19 cases, it is important to introduce an automated detection system as an alternative diagnostic method to prevent the spread of COVID-19 [7].

Meanwhile, Gozes et al. [40] employed a deep-learning approach to automatically identify COVID-19 patients and examine the disease burden quantification on CT scans using a dataset of CT scans from 157 foreign patients from China and the USA. Their proposed system analyses the CT scan at two distinct levels: subsystems A and B. Subsystem A performs a 3D analysis, and subsystem B performs a 2D analysis of each segment of the scan to identify and locate broader diffuse opacities, including ground-glass infiltrates (which have been clinically identified as representative of COVID-19). To evaluate their system, the authors applied Resnet-50-2 to subsystem B and obtained an area under the curve of 99.6%. The sensitivity and specificity were 98.2% and 92.2%, respectively.

Moreover, Wang et al. [38] developed a deep-learning approach for extracting information from CT scans. Their study included a collection of 453 CT scans from 99 patients. They extracted 195 regions of interest (ROIs) of sizes ranging from 395×223 to 636×533 pixels from the CT scans of 44 COVID-19-positive pneumonia patients and 258 ROIs from those of 50 COVID-19-negative patients. They applied a modified network inception model and obtained an accuracy of 82.9% for the internal validation with a specificity of 80.5% and a sensitivity of 84%. The external testing dataset exhibited a total accuracy of 73.1% with a specificity of 67% and a sensitivity of 74%.

Fu et al. [41] proposed a classification system based on ResNet-50 to detect COVID-19 and some other infectious lung diseases (bacterial pneumonia and pulmonary tuberculosis). The authors collected a dataset of 60,427 CT scans from 918 patients; 14,944 of these CT scans were from 150 COVID-19 patients and 15,133 from 154 non-COVID-19 viral pneumonia patients. They performed several tests for several lung diseases. The achieved accuracy, sensitivity, and specificity were 98.8%, 98.2%, and 98.9%, respectively.

Xu et al. [42] reported that real-time RT-PCR has a low positive rate at the early stage of COVID-19. They developed an early screening model that uses deep-learning techniques for distinguishing COVID-19 pneumonia from influenza (a viral pneumonia) and stable cases using pulmonary CT images. A dataset of 618 CT samples was obtained for the analysis, and the images were classified as COVID-19, influenza (a viral pneumonia), and other cases using ResNet-18 and ResNet-based methods. The authors employed a noisy or Bayesian function to differentiate the infected images and obtained a detection accuracy of 86.7%.

## 2) COVID-19 Infection Prediction Using Machine Learning Techniques

ML is the science of training machines using mathematical models to learn and analyse data. Once ML is implemented in a system, the data are analysed, and interesting patterns are detected. The validation data are then categorised according to the patterns learned during the learning process. As COVID-19 infection has rapidly spread worldwide and international action is required, it is important to develop a strategy to estimate the number of potentially infected people on a regular basis to adopt the appropriate measures. Currently, decision-makers rely on certain decision-making statistics such as imposing lockdowns on infected cities or countries. Therefore, ML can be used to predict the behaviours of new cases to stop the disease from spreading.

Li et al. [43] developed a prediction model using ML algorithms to combat COVID-19 in mainland China and in other infected countries in the world. The authors developed a model to estimate the number of reported cases and deaths in mainland China and in the world. The data used to build the models were collected between 20 January 2020 and 1 March 2020. They estimated that the peak of the COVID-19 outbreak in mainland China occurred on 22 February 2020 and on 10 April 2020 worldwide. The authors also stated that COVID-19 would be controlled at the beginning of April 2020 in mainland China and in mid-June 2020 across the world. They concluded that the estimated number of COVID-19 cases would be approximately 89,000 in China and 403,000 worldwide during the outbreak. As of 17 April 2020, the estimated number of deaths was 4000 in mainland China and 18,300 worldwide. It is clear that their forecast was similar to the actual situation in China as the total numbers of infected cases and deaths had exceeded 82,367 and 3342, respectively. However, the number of confirmed cases worldwide exceeded their estimations as the numbers of infected cases and deaths exceeded 2 million and reached 145,416 as of 17 April 2020, respectively [2].

Kumar et al. [44] predicted the COVID-19 spread in the 15 most-infected countries in the world using the autoregressive integrated moving average (ARIMA) model. The outcome of their prediction indicates that circumstances would worsen in Iran and Europe, especially in Italy, Spain, and France. Moreover, their prediction indicated that the number of cases in South Korea and mainland China would become more stable. The study also indicated that COVID-19 would spread exponentially in the USA and that strict official measures

would be urgently required to stop the disease from spreading. Although the prediction of COVID-19 cases for the USA was 1 million between 8 April and 30 April 2020, it reached 677,570 on 17 April 2020. Furthermore, Italy had 168,941 cases, although it was predicted to have 300,000 cases [2].

Huang et al. [45] applied a CNN to a limited dataset, which was not specifically defined in their study, to evaluate and estimate the number of reported cases in China. The authors used the mean absolute and root mean square errors to compare their model with other deep-learning models, including multilayer perceptron, long short-term memory (LSTM), and gated recurrent units. The authors concluded that the obtained results promise a high predictive efficiency.

Pandey et al. [46] utilised two statistical algorithms—the susceptible-exposed-infectious-recovered (SEIR) and regression models—to evaluate and forecast the distribution of COVID-19 in India. They used a dataset retrieved from the John Hopkins University repository. The prediction results from the SEIR and regression models showed that the number of confirmed COVID-19 cases would reach 5300 and 6153 cases, respectively, by 13 April 2020. However, the confirmed cases in India on that date were 10,453 and 6153 for the SEIR and regression models, respectively [2].

### III. System Design

Our proposed deep learning-based COVID-19 detection comprises several phases, as illustrated in Figure 1. The phases are summarised in the following five steps:

**Step 1:** Collect the chest X-ray images for the dataset from COVID-19 patients and healthy persons.

**Step 2:** Generate 1000 chest X-ray images using data augmentation.

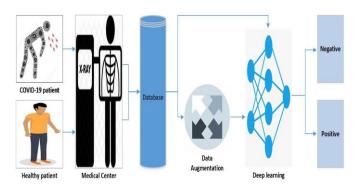
**Step 3:** Represent the images in a feature space and apply deep learning.

**Step 4:** Split the dataset into two sets: a training set and a validation set.

**Step 5:** Evaluate the performance of the detector on the validation dataset.

#### A. Dataset

Two types of datasets were used in the evaluation, the original dataset (without augmentation) and the augmented dataset, which are summarised in Tables 1 and 2, respectively. The dataset contained the following: a) a healthy dataset containing chest X-ray images of healthy persons and b) a COVID-19 dataset containing chest X-ray images of COVID-19 patients. The original dataset was obtained from the Kaggle database, and its total number of images is 128, as presented in Table 1 [47].



**Figure 1**. Architecture of the proposed system

Owing to the limited availability of chest X-ray images, we generated our dataset using data augmentation [48]. Data augmentation is an AI method for increasing the size and the diversity of labelled training sets by generating different iterations of the samples in a dataset. Data augmentation methods are commonly used in ML to address class imbalance problems, reduce overfitting in deep learning, and improve convergence, which ultimately contributes to better results. The total number of images in the dataset became 1000 after applying augmentation, as presented in Table 2 [47].

Table 1 Original dataset (without augmentation)

X-ray images	Number
Healthy	28
COVID-19	70
Total	128

Table 2 Augmented dataset

X-ray images	Number
Healthy	500
COVID-19	500
Total	1000

#### B. Environment

A computer with Microsoft Windows 10 was used for the experiment. It has the following specifications: Intel Core i7-8565U 1.80-GHz processor, 16 GB of DDR4 RAM, and 1 TB of hard disk. We installed the virtual machine tool VMware Workstation Pro version 14.1.8 build-14921873 on it. Then, we installed Ubuntu 18.04.4 (64 bit) on the virtual machine and the following libraries and software:

• ARIMA:

 $\frac{https://www.statsmodels.org/stable/generated/statsmodels.ts}{a.arima\_model.ARIMA.html}$ 

- Fbprophet: <a href="https://pypi.org/project/fbprophet/">https://pypi.org/project/fbprophet/</a>
- ImageDataGenerator: <a href="https://keras.io/preprocessing/image/">https://keras.io/preprocessing/image/</a>
- Keras: https://keras.io/
- •LSTM:

 $\frac{https://www.tensorflow.org/api\ docs/python/tf/keras/layers/LS}{TM}$ 

- Matplotlib: <a href="https://matplotlib.org/">https://matplotlib.org/</a>
- NumPy: <a href="https://numpy.org/">https://numpy.org/</a>
- Pandas: <a href="https://pandas.pydata.org/">https://pandas.pydata.org/</a>
- Python: <a href="https://www.python.org/">https://www.python.org/</a>
- Scikit: https://scikit-learn.org/
- SciPy: <a href="https://www.scipy.org/">https://www.scipy.org/</a>
- TensorFlow: <a href="https://www.tensorflow.org/">https://www.tensorflow.org/</a>

All the results and predictions made in this study have been uploaded to the Kaggle database [49, 50]. We believe that by making the system and solution publicly available, we draw attention to the most affected areas, thereby preventing the spread of the COVID-19 outbreak and fostering the use of deep-learning techniques in COVID-19 research.

#### C. Evaluation Metrics

To assess the reliability of the proposed deep learning-based COVID-19 detector, we adopted the same metrics as those used by Alazab et al. [51-54] and considered the following standard metrics: precision, recall, and F-measure. These metrics are calculated on the basis of the true-positive (TP),

true-negative (TN), false-positive (FP), and false-negative (FN) scores:

- TP is the proportion of positive COVID-19 chest X-ray images that were correctly labelled as positive.
- FP is the proportion of negative (healthy) COVID-19 chest X-ray images that were mislabelled as positive.
- TN is the proportion of negative (healthy) chest X-ray images that were correctly labelled as healthy.
- FN is the proportion of positive COVID-19 chest X-ray images that were mislabelled as negative (healthy).
- Accuracy: This metric measures the percentage of correctly identified cases relative to the entire dataset. The ML algorithm performs better if the accuracy is higher. Accuracy is a significant measure for a test dataset that includes a balanced class. It is computed as follows:

$$Accuracy = (TP + TN)/(TP + TN + FP + FN)$$
(1)

 Precision: This metric is a measure of exactness, which is calculated as the percentage of positive predictions of COVID-19 that were true positives divided by the number of predicted positives. It is computed as follows:

$$Precision = TP/(TP + FP)$$
 (2)

 Recall: This metric is a measure of completeness, which is calculated as the percentage of positives that were correctly identified as true positives divided by the number of actual positives. It is computed as follows:

$$Recall = TP/(TP + FN)$$
 (3)

 F-measure: This is a combination of precision and recall that provides a significant measure for a test dataset that includes an imbalanced class. It is computed as follows:

$$F-Measure = 2 \left( \frac{Precision \times Recall}{(Precision + Recall)} \right)$$
 (4)  
Root Mean Square Error (RMSE): This metric

■ Root Mean Square Error (RMSE): This metric measures the differences between the actual  $(x_i)$  and the predicted  $(\hat{x}_i)$  numbers of COVID-19 confirmations, recoveries, and deaths (N). The main advantage of RMSE is that it penalises large prediction errors. RMSE was used to compare the prediction errors of the three prediction algorithms. It is defined as follows:

**RMSE** = 
$$\sqrt{\frac{1}{N}} \sum_{i=1}^{N} (x_i - \hat{x}_i)^2$$
 (5)

 Correlation coefficient: This metric is often used to evaluate the performance of a prediction algorithm. It is defined as follows:

$$CC = \left(1 - \frac{1}{N} \sum_{i=1}^{N} |x_i - \hat{x}_i|\right) * 100\%$$
 (6)

#### **IV.** Experimental Results

Firstly, we examined the most infected areas across the world. In Section 4.1, we show that coastal areas are heavily affected by the COVID-19 outbreak as the number of cases in those

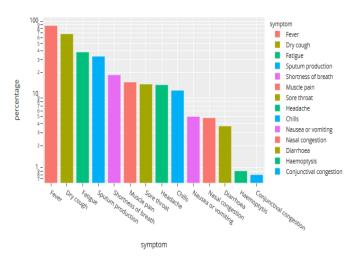
areas is significantly higher than those in other non-coastal areas (Figures. 3–6).

Secondly, we predicted the number of COVID-19 confirmations, recoveries, and deaths in Jordan and Australia over the next 7 days using three well-known time-series forecasting algorithms: PA (prophet algorithm), ARIMA, and LSTM. In Section 4.2, we show how the application of these algorithms allowed us to estimate the forecasting outcomes for certain countries with a detection rate of 99% (Tables 3 and 4) and (Figures. 7 -13).

Thirdly, we examined whether chest X-ray images can be used to develop sophisticated classification models for COVID-19 prediction. In Section 4.3, we present the application of a deep-learning algorithm on two datasets. Empirical findings indicated that our proposed system is reliable in detecting COVID-19 and has an F-measure range of 95–99%, as revealed by Figures. 15 -16.

#### A. Coronavirus Statistics

SARS-CoV-2 is a new family of viruses that has never been encountered before. The virus was first discovered in pangolins before its spread to humans [55]. The typical symptoms of COVID-19 include fever, dry cough, fatigue, sputum production, shortness of breath, sore throat, headache, chills, nausea or vomiting, nasal congestion, diarrhoea, haemoptysis, and conjunctival discomfort, although some patients also suffer from general tiredness, runny nose, and loss of taste and/or scent. Figure 2 shows a bar graph of the common COVID-19 symptoms sorted by their percentage of occurrences [56].



**Figure 2**. Common COVID-19 symptoms

In this section, the connection between coastal and non-coastal areas is further explored as a large proportion of infected cases were recorded in coastal areas. On the basis of the collected statistics, COVID-19 has a rapid spread in coastal areas. The following examples support this conclusion:

In Australia, there were more than 7000 confirmed cases. The highest number of confirmed cases were in New South Wales, Victoria, Queensland, Western Australia, South Australia, Tasmania, and Australian Capital Territory and Northern Territory [57]. Figure 3 reveals the most affected areas in Australia.



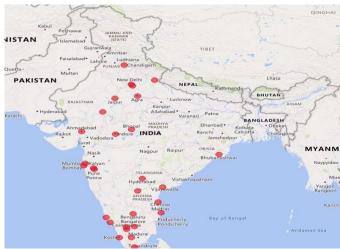
**Figure 3.** Areas with the highest number of confirmed cases in Australia

• In South Korea, the disease peaked on 20 January 2020, and the number of confirmed cases exceeded 11,000. Figure 4 shows that the regions that were most severely affected by COVID-19 were coastal or near-coastal areas.



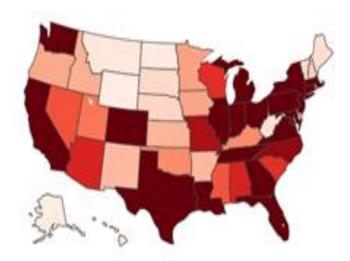
Figure 4. Areas with the highest number of confirmed cases in South Korea

• In India, the number of reported confirmed cases was more than 158,000, and the most-affected places were Maharashtra; Tamil Nadu, Delhi, Gujrat, Rajasthan, Madhya Pradesh, Uttar Pradesh etc. as shown in Figure 5 [58].



**Figure 5.** Areas with the highest number of confirmed cases in India

• In the USA, there were more than 1,745,843 confirmed cases on 27 May 2020. The first case was found in Oregon, which is located in the Pacific Coast. Coastal states, including Washington, Oregon, California, Arizona, and Texas, reported high numbers of confirmed cases. Furthermore, states including Wisconsin and Illinois with long lake coastlines also reported confirmed cases at the initial stage of the COVID-19 spread. Other eastern coastal states including New York, Maine, New Hampshire, Massachusetts, Rhode Island, Connecticut, New Jersey, Delaware, Maryland, Virginia, North Carolina, South Carolina, Georgia, Florida, and Indiana also reported high numbers of confirmed cases, as well as other coastal areas such as Colorado and Nebraska. Thus, most of the states that reported the highest numbers of cases are located in the coastal regions. Figure 6 highlights the coastal regions with the highest number of cases in maroon.



**Figure 6.** Areas with the highest number of confirmed cases in USA

#### B. Prediction

In our system, we predicted the number of COVID-19 confirmations, recoveries, and deaths in Jordan and Australia using the following three well-known forecasting algorithms: i) PA [59], ii) ARIMA [60], and iii) LSTM [61]. These algorithms were trained to make predictions for the next 7 days from datasets that were collected from a statistics website [2].

Originally proposed by Taylor and Letham, PA is one of the well-known algorithms for solving multi-seasonal timeseries forecasting problems on Facebook [59]. It additively decomposes a time series-based forecasting problem into trends, seasonal, remainder, and holiday components. It achieves time-series forecasting using the following relation:

$$x_{t} = \hat{S}_{t}^{1} + \hat{S}_{t}^{2} + \dots + \hat{S}_{t}^{n} + \hat{T}_{t} + \hat{R}_{t} + \hat{H}_{t}$$
 (7)

 $x_t = \hat{S}_t^1 + \hat{S}_t^2 + \dots + \hat{S}_t^n + \hat{T}_t + \hat{R}_t + \hat{H}_t \qquad (7)$  where  $x_t$  represents an observation at time t, n denotes the number of distinct seasonal patterns,  $\hat{S}_t^n$  represents the seasonal effect,  $\hat{T}_t$  represents the effect of trend,  $\hat{R}_t$  represents the remainder component of the observation and  $\hat{H}_t$  denotes the holiday covariance, which represents the effect of holidays.

ARIMA is a simplified type of autoregressive moving average model that incorporates autoregressive and movingaverage models to develop a composite forecasting model. The autoregressive model utilises the dependency between the observations and several lagged observations, whereas the moving-average model uses the association between the

observations and the residual error values by using the moving average for the lagged observations. ARIMA uses the order factors p, d, and q, where p is the order of the autoregressive model, d is the order of the differencing, and q is the order of the moving average. The algorithm is computed as follows:

ing average. The algorithm is computed as follows:
$$y = c + \epsilon_t + \sum_{l=1}^{p} \emptyset_l y_{t-l}$$

$$+ y = c + \epsilon_t + \sum_{l=1}^{q} \theta_l \epsilon_{t-1}$$
is an independent and homogeneously dist

where  $\epsilon_t$  is an independent and homogeneously distributed error term, c is a constant term, y is an actual value at time t, and  $\phi$  and  $\theta$  are the tuning parameters of the autoregressive and moving-average models, respectively.

LSTM is a form of a recurrent neural network (RNN) that memorises earlier patterns in data sequences. It was originally proposed by Hochreiter and Schmidhuber [62]. It replaces the hidden layer neurons of the RNN with a series of memory cells. The key is the state of the memory cell that filters data using a gate structure that updates the state of the memory cell. It includes the input, forgotten, and output gates for its gate structure. Each cell has three sigmoid layers and one tanh layer, as shown in Figure 7 [61].

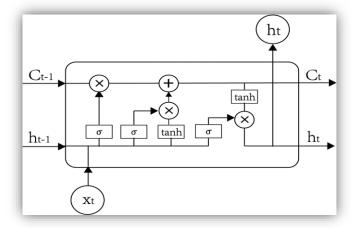


Figure 7. Structure of a memory cell in an LSTM

(i) The forgotten gate determines the cell state data that must be discarded. Each memory cell of the LSTM combines the previous output  $h_{t-1}$  at time t-1 and the external current data  $x_t$  in a vector  $[h_{t-1}, x_t]$  through a special sigmoid function  $\sigma$ , shown as follows:

$$f_t = \sigma(W_f.[h_{t-1}, x_t] + b_f)$$
 (9)

where  $W_f$  and  $b_f$  are the weight and the bias, respectively, of the forgotten gate. On the basis of its inputs, if the forgotten gate outputs '1', it indicates 'reserve', and if it outputs '0', it indicates 'discard' data in a cell.

(ii) The input gate determines the period necessary to preserve the current data  $x_t$  in a cell  $C_t$ . It finds the state of the cell  $C_t$ , and the data value is updated by the sigmoid layer, shown as follows:

$$i_t = \sigma(W_f. [h_{t-1}, x_t] + b_f)$$
 (10)

The input gate updates the data that need to be upgraded to cell  $C_t$ . A new vector  $\hat{C}_t$  is thus created by the tanh layer to determine the amount of data that must be added, and it is defined as follows:

$$\hat{C}_t = tanh(W_c.[h_{t-1}, x_t] + b_c)$$
 (11)

Finally, the state of the memory cell is updated as follows:

$$C_t = f_t * C_{t-1} + i_t * \hat{C}_t \tag{12}$$

(iii) The output gate controls how much of the current cell state is discarded. The data are determined by a sigmoid layer. The cell state is processed by the tanh layer and multiplied by the output retrieved from the sigmoid layer to obtain the final output of the cell, as shown below:

 $h_t = \sigma(W_\sigma.[h_{t-1},x_t] + b_o) * \tanh(C_t)$  (13) To assess the performance of the implemented forecasting algorithms, we collected the numbers of COVID-19 confirmations, recoveries, and deaths between 30 March 2020 and 5 April 2020 in Australia and Jordan. The predicted values were then compared with the real numbers, and the results are presented in figures. 8–13. Figures 8–10 show the predicted COVID-19 confirmations, recoveries, and deaths in Australia from 30 March to 5 April 2020 using the PA, ARIMA, and LSTM algorithms, respectively. Figures 11–13 show the predicted COVID-19 confirmations, recoveries, and deaths in Jordan from 30 March 2020 to 5 April 2020. The three algorithms were compared in terms of RMSE, accuracy, and correlation coefficient, as detailed in Tables 3 and 4.

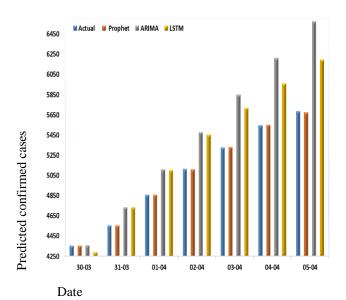


Figure 8. Predicted COVID-19 confirmed cases in Australia

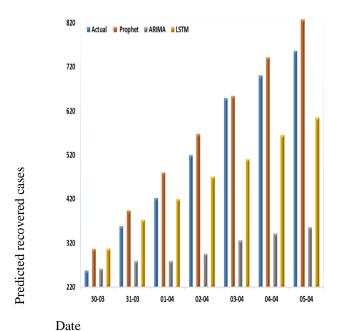


Figure 9. Predicted COVID-19 recovered cases in Australia

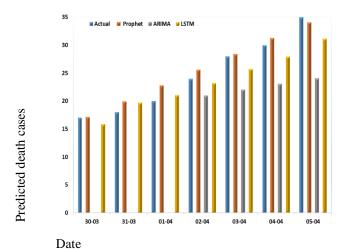
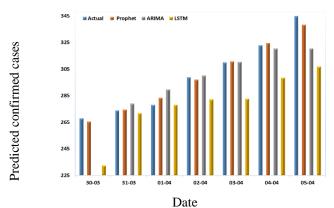


Figure 10. Predicted COVID-19 death cases in Australia



**Figure 11.** Predicted COVID-19 confirmed cases in Jordan

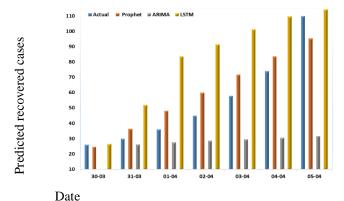


Figure 12. Predicted COVID-19 recovered cases in Jordan

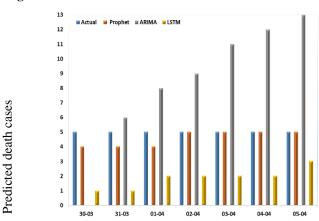


Figure 13. Predicted COVID-19 death cases in Jordan

Table 3 details the performances of the three prediction algorithms in Australia. We calculated the correlation coefficient, accuracy, and RMSE for each algorithm. PA delivered the best results, whereas ARIMA delivered the worst results. Moreover, PA predicted the numbers of COVID-19 confirmations, recoveries, and deaths and obtained prediction accuracies of 99.94%, 90.29% 94.18%, respectively.

Table 4 details the performances of the three prediction algorithms in Jordan. We calculated the correlation coefficient, accuracy, and RMSE for each algorithm. PA delivered the best results, whereas LSTM delivered the worst results. Moreover, PA predicted the numbers of COVID-19 confirmations, recoveries, and deaths and obtained prediction accuracies of 99.08%, 79.39%, and 86.82%, respectively.

Table 3 Results of three prediction algorithms in Australia

Prediction	Correlation	Accuracy	RMSE
Algorithm	Coefficient		
PA confirmed cases	0.99	99.94%	03.94
PA (recovered cases)	0.99	90.29%	47.83
PA (death cases)	0.98	94.18%	01.55
ARIMA (confirmed	0.99	92.33%	497.55
cases)			
ARIMA (recovered	0.98	63.52%	260.12
cases)			
ARIMA (death	0.98	78.02%	07.26
cases)			
LSTM (confirmed	0.99	94.16%	337.18
cases)			
LSTM (recovered	0.99	86.44%	97.36
cases)			
LSTM (death cases)	0.98	92.76%	02.07

Table 4 Results of the three prediction algorithms in Jordan

Prediction	Correlation	Accuracy	RMSE
Algorithm	Coefficient		
PA (confirmed	0.99	99.08%	03.51
cases)			
PA (recovered	0.94	79.39%	11.42
cases)			
PA (death cases)	-	86.82%	00.78
ARIMA (confirmed	0.95	97.59%	11.30
cases)			
ARIMA (recovered	0.93	57.79%	39.26
cases)			
ARIMA (death		12.87 %	05.59
cases)			
LSTM (confirmed	0.85	93.23%	25.02
cases)			
LSTM (recovered	0.94	37.33%	11.42
cases)			
LSTM (death cases)		39.97%	03.04

#### C. Deep Learning

This study developed a CNN-based COVID-19 detection model that was tested with both the original and the augmented datasets. All the chest X-ray images used were resized to 224×224 pixels while ignoring the aspect ratio. Figures 14a and b present the chest X-ray images of healthy and COVID-19-infected patients, respectively. The collected dataset was randomly split into a training data subset and a testing data subset.





(a) Healthy Figure 14. Chest X-ray images

(b) Coronavirus Infected

The COVID-19 detection model is based on VGG16, a type of CNN, and it is designed to detect COVID-19 using chest X-ray images. VGG16 is one of the highest-quality visionmodel architectures currently available. It generates evidence in favour of stacking convolutional layers with tiny filters  $(3\times3)$  rather than using a single layer with larger filter sizes  $(5\times5 \text{ and } 7\times7)$  as stacked  $3\times3$  filters will approximate bigger ones [63].

The COVID-19 detector was trained and tested on the collected dataset, 80% of which was used for training and the remaining 20% was used for testing. The weights of the CNN were randomly initialised, and the batch size was varied up to 25 and empirically set to 25 to avoid overfitting and to achieve the highest training accuracy. Furthermore, the learning rate was initially set to 0.1. Figure 15 details the accuracy of the COVID-19 detector and its loss values for the implemented detector with augmentation. Figure 16 details the accuracy of the same detector with augmentation and its loss values.

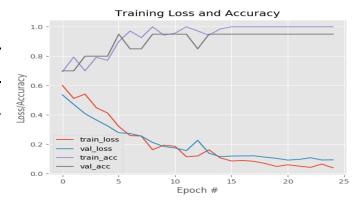


Figure 15. Performance of CNN models in the original dataset

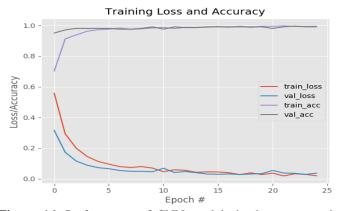
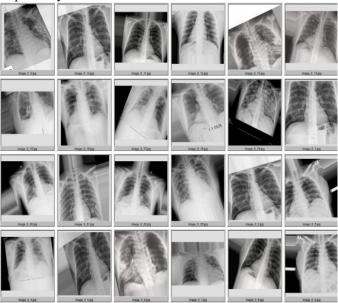
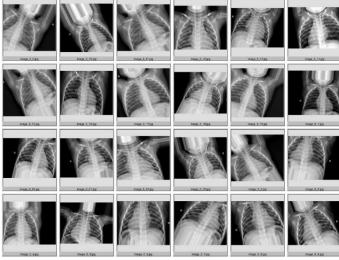


Figure 16. Performance of CNN models in the augmented dataset

Special attention must be paid to the avoidance of overfitting in the un-augmented dataset, especially when increasing the epochs as the validation slowly improves in the beginning and then stops improving when the epochs are increased, as shown in Figure 15. When the augmented dataset is used, the gap between the training and validation becomes smaller after a few epochs, as shown in Figure 16. Thus, a greater improvement is achieved in the training process, and a more generalised and robust COVID-19 detector is achieved using the CNN models when implementing data augmentation on the dataset. Figures 17 and 18 show the augmented chest X-ray images of COVID-19 patients and healthy persons, respectively.



**Figure 17.** Augmented chest X-ray images for COVID-19 patients



**Figure 18.** Augmented chest X-ray images for healthy people

The CNN-based COVID-19 detector trained on an unaugmented dataset achieved a weighted average F-measure of 95%. The same COVID-19 detector achieved a weighted average F-measure of 99% when trained on an augmented dataset, as shown in Figure 19. Hence, the COVID-19 detector exhibits superior performance metrics in terms of recall, precision, and F-measure when trained on augmented data. It is therefore sufficiently robust and helpful for rapidly diagnosing a large number of suspected COVID-19 patients.

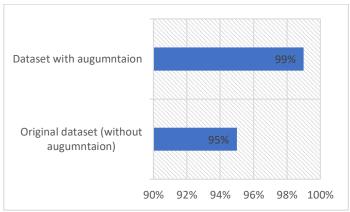


Figure 19. F-measure scores

#### V. Discussions

This study provided a forecasting analysis of COVID-19 confirmations, recoveries, and deaths in Australia and Jordan. It further implemented a CNN-based COVID-19 detector to identify COVID-19 infections using X-ray images. Based on the study results, the following conclusions were drawn:

- PA delivered the best performance for COVID-19 prediction over 7 days, compared to LSTM and ARIMA.
- The predictions will enable people in both countries to predict their medical needs for tackling the spread of COVID-19.
- ARIMA cannot make predictions over the next 1, 2, and 3 days.
- After investigating the number of COVID-19 confirmations, recoveries, and deaths in various countries, we found that coastal areas are significantly impacted by the disease because the numbers of cases in those areas are significantly higher than those in other non-coastal areas. This observation is medically consistent with the propagation capability of viruses in areas with higher humidity rates. Thus, the authors advise healthcare professionals to devote greater attention to coastal regions.
- The use of chest X-ray images is recommended for diagnosing COVID-19 because X-rays are easily obtained at nearby hospitals or clinics fairly quickly and at low costs.
- Our CNN-based COVID-19 detector delivered superior performance in terms of precision, recall, and F-measure.
- The application of ML techniques for COVID-19 diagnosis using our CNN-based COVID-19 detector is recommended.
- It is well known that VGG16 (Wu et al., 2017) outperforms many convolutional networks, such as GoogLeNet and SqueezeNet, and its feature representation capability is beneficial for classification accuracy. Hence, VGG16 is a recommended version of a deep CNN-based algorithm as it makes training easier and quicker. It was implemented in our COVID-19 detector to improve its accuracy in diagnosing COVID-19 in chest X-ray images.

Our COVID-19 detector obtained better results when using augmentation. A better training process was achieved as the gap between the training and validation became smaller. Moreover, a more generalized and robust COVID-19 detector was achieved as the F-measure improved from 0.95 to 0.99. Thus, the COVID-19 detector trained on augmented data provides superior performance metrics and is robust for diagnosing COVID-19 in chest X-ray images.

#### VI. Conclusions and Future Work

The rapid spread of COVID-19 across the world and the increasing number of deaths require urgent actions from all sectors. Future prediction of potential infections will enable authorities to tackle the consequences effectively. Furthermore, it is necessary to keep up with the number of infected people by performing regular check-ups, and it is often vital to quarantine infected people and adopt medical measures. Additionally, attention should be given to several other factors to curb the spread of COVID-19, such as the environmental effects and the similarities among the most affected areas, and careful measures should be adopted. In this paper, AI-based techniques were proposed for the prediction and diagnosis of COVID-19:

- Prediction models such as the PA, ARIMA, and LSTM algorithms were used to predict the number of COVID-19 confirmations, recoveries, and deaths over the next 7 days. PA delivered the best performance. It predicted the number of COVID-19 confirmations, recoveries, and deaths in Australia and obtained prediction accuracies of 99.94%, 90.29%, and 94.18%, respectively. It also predicted the number of COVID-19 confirmations, recoveries, and deaths in Jordan and obtained prediction accuracies of 99.08%, 79.39%, and 86.82%, respectively. Meanwhile, investigation into more sophisticated forecasting and prediction methods is a subject of a future work.
- A diagnosis model using VGG16 was proposed to detect COVID-19 using chest X-ray images. The model allows the rapid and reliable detection of COVID-19, enabling it to achieve an F-measure of 99% using an augmented dataset. In a future study, we will consider diagnosing COVID-19 in chest CT scan images using the VGG-XX versions and compare their performances using larger datasets.

A further contribution of this study is the analysis of the COVID-19 spread and its related statistical data based on its global regional distributions. Thus, two main conclusions were drawn using our AI-based analysis: (i) the most highly infected areas have similar characteristics, and (ii) the spread of the disease in coastal areas is significantly higher than that in other non-coastal areas. Therefore, extra care and attention should be given to coastal cities. In our future work, we will investigate the effects of temperature, humidity, and terrain on the COVID-19 spread in cities and countries.

#### Acknowledgment

This work was supported by the faculty of Artificial intelligence, Al-Balqa Applied University.

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