

# Predicting Covid-19 Cases using CNN Model

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**Abstract:** The prediction of COVID-19 confirmed cases is a complex time-series problem. In the literature, Long Short Time Memory (LSTM) has proven its efficiency to resolve issues related to the time series problems. However, Convolution neural network (CNN) did not been widely used in this aim and is considered as more suitable for imaging processing. Therefore, in this paper, we use it to predict COVID-19 cases and compared it with LSTM in the context of Morocco during the period of confinement. The obtained results which we present and discuss in this article are very promising.

## 1 INTRODUCTION

Reported first in December 2019, in Wuhan, China, Coronavirus 2019 (COVID-19), caused by the strain of coronavirus named “Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2)”, has become now, according to the World Health Organization WHO, an international epidemic having caused 17 660 523 infected cases, and 680 894 deaths on August 03, 2020 (Organization, 2020).

Since the first cases were reported, multiple academic and medical works have explored various approaches to find a solution for the COVID-19 epidemic in different search areas including the Machine Learning (ML). Some ML, and especially Deep Learning (DL), work interested in COVID-19 are oriented to help medical staff to efficiently diagnostic infected people (Singh, Kumar, Vaishali, & Kaur, Coronavirus (COVID-19) Classification using CT Images by Machine Learning Methods, 2020) (Wang, et al., 2020) (Gozes, et al., 2020) (Metsky, Freije, Kosoko-Thoroddsen, & Myhrvold, 2020), and others are directed to analyze the pandemic situation like predicting new patients to contribute to local hospital arrangement (Alimadadi, et al., 2020) (Bouhamed, 2020) (Chimmula & Zhang, 2020) (Pinter, Felde, MOSAVI, Ghamisi, & Gloaguen, 2020).

In general, a lot of work applied ML and DL to predict COVID-19 new cases use time-series

forecasting models (Chimmula & Zhang, 2020) (Zeroual, Harrou, Dairid, & Sun, 2020) (Azarafza, Azarafza, & Tanha, 2020) (Punn, Sonbhadra, & Agarwal, 2020). Actually, Recurrent Neural Networks (RNN) architecture are adopted by several researchers thanks to their capacity in handling time-dependent datasets, and hence they were widely used in the context of predicting COVID-19 new cases. Moreover, other Neural Networks (NN) architectures were used for this objective.

The objective of this work is to use Convolution Neural Network (CNN) architecture in time-series forecasting to predict the COVID-19 new cases in Morocco. We compare our results with a Long Short-Term Memory (LSTM) neural network explored in the same context by another work conducted by our research team.

This paper is constituted of five sections. The second one explores the related work, which is relative to CNN architectures and their use in Time-series forecasting, especially in the context of COVID-19. The third section is about Materials and Method used to perform our experimental studies. We present and discuss the results in the fourth section, and finally we conclude in the fifth section.

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## 2 RELATED WORK

Most often, a time-series is a succession of points equidistant in time. It can be thought of as sequence of vectors  $s(t)$ , with  $t$  between 0 to  $n$ , where  $t$  denotes the spent time,  $x$  is a value varying with  $t$ , and  $n$  is the max value of  $t$ . The problem of forecasting future values of  $s$  from values of  $s$  up to the current time can be stated as finding a function  $f: \mathbb{R}^N \rightarrow \mathbb{R}$  such as to estimate  $s$  at time  $t + d$ , from the  $N$  time steps back from time  $t$  (Frank, Davey, & Hunt, 2001):

$$s(t + d) = f(s(t), s(t - 1), \dots, s(t - N + 1)) \quad (1)$$

In the literature, RNN architectures were considered the most suitable for solving problems that involve sequential data (Ganatra & Patel, 2018).

LSTM is a special type of recurrent neural network with long and short term memory cells (Kamal, Bae, Sunghyun, & Yun, 2020). It was proven that it has good accuracy in the context of COVID-19 prediction. In (Azarafza, Azarafza, & Tanha, 2020), LSTM was more efficient than RNN in predicting infections in Iran, while LSTM learning has given good RMSE and suggested that the epidemic's expected outcome will be reached in the period of June 2020 in (Chimmula & Zhang, 2020) but the actual situation proves that it was not true. We think it is because of the small size of the analyzed data set (confirmed cases until March 31, 2020). Nevertheless, comparing LSTM, RNN, GRU, and Variational AutoEncoder VAE have proven the efficiency of VAE (Zeroual, Harrou, Dairid, & Sun, 2020).

The CNN architectures as shown in Figure 1, in the other side, were basically designed to extract local relationships that are invariant across spatial dimensions. Commonly they have three types of layers; including fully-connected layer, pooling layer, convolutional and non-linearity layers (Abiodunab, et al., 2018).

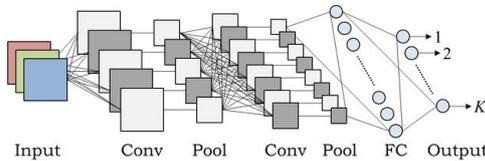


Figure 1: An example of CNN architecture (Hidaka & Kurita, 2017).

To adapt CNNs to time-series datasets, researchers use multiple layers of causal convolutions (Oord, et al., 2016) (Bai, Kolter, & Koltun, 2018).

For an intermediate feature at hidden layer  $l$ , each causal convolutional filter takes the form below (Lim & Zohren, 20):

$$h_l^{t+1} = A((W * h)(l, t)) \quad (2)$$

$$(W * h)(l, t) = \sum_{\tau=1}^k w(l, \tau) h_{t-\tau}^l \quad (3)$$

where  $h_l^{t+1} \in \mathbb{R}^{H_{in}}$   $H_{in}$  is an intermediate state at layer  $l$  at time  $t$ ,  $*$  is the convolution operator,  $W(l, \tau) \in \mathbb{R}^{H_{out} \times H_{in}}$  is a fixed filter weight at layer  $l$ , and  $A(\cdot)$  is an activation function representing any architecture-specific non-processing. In practice, the one-dimensional 1-D CNN architecture is used to classify and predict future time-series values.

In the context of COVID-19 analysis, the CNN architectures were widely used to detect it from X-ray or Computed Tomography CT images. On the X-ray images side, authors in (Apostolopoulos & Mpesiana, 2020) utilized a dataset of X-beam pictures from patients with regular bacterial pneumonia, affirmed Covid-19 sickness, and ordinary occurrences to distinguish the COVID-19 infection. Similarly, in (Ozturk, et al., 2020), the authors use a dataset of raw chest X-ray images with the DarkNet model (based on CNN architecture) to give exact diagnostics, classify people with the COVID for those who don't have the COVID-19 and for multi-class classification between COVID contaminated people, healthy individuals and persons with pneumonia. Authors in (Minaee, Kafieh, Sonka, Yazdani, & Soufie, 2020) used a transfer learning on a subset of 2000 x-ray images to learn four successful CNNs models, comprising ResNet18, ResNet50, SqueezeNet, and DenseNet-121, to recognize COVID-19 illness in a chest image dataset collected from the publicly available datasets, and images exhibiting COVID-19 disease presence were identified by board-certified radiologist. While authors in (Islam & Asraf, 2020) use a dataset of 4575 X-ray images, including 1525 images of COVID-19, the CNN to extract for deep feature and LSTM to detect COVID-19 using the extracted feature. On the other side, authors (Singh, Kumar, Vaishali, & Kaur, Classification of COVID-19 patients from chest CT images using multi-objective differential evolution-based convolutional neural networks, 2020) use a dataset of chest CT images and a CNN model to assign COVID-19 infected patients as infected and uninfected in another case. They calibrated the settings of CNN based on the multi-objective differential evolution. Similarly, authors in (Wang, et al., 2020) use a dataset of 1,065 CT images of pathogen-confirmed COVID-19 cases

(325 images) along with those previously diagnosed with typical viral pneumonia (740 images) and a transfer-learning model to train a CNN model followed by an internal and an external validation.

The CNN did not have the same popularity in the context of time-series forecasting of COVID-19 cases. Up to day, there are only four works that have exploited the CNN architectures to predict COVID-19, in the literature. The first one is (Amo-Boateng, 2020) where 1D-CNN is used on the chronological data records of known COVID-19 positive cases in order to predict the steps for reporting countries and territories. The second one is (Barman, 2020) that explores the performance of Auto-Regressive Integrated Moving Average (ARIMA) model and several LSTM models, including the CNN-LSTM, in forecasting the number of confirmed COVID-19 cases. According to the authors, ARIMA marked better results. The third one is (Dutta, Bandyopadhyay, & Kim, 2020) a study that investigates the forecast of the growth of COVID-19 at a close later time in the world through the CNN and CNN-LSTM models. The results demonstrate that the mixed CNN-LSTM framework is more efficient than the rest of the models. The last one is (Huang, Chen, Ma, & Kuo, 2020) that proposes a CNN model to analysed and predict the number of confirmed cases in several cities with the most confirmed cases in China. The findings of the research stated that, in comparison to a number of alternative deep learning methods (Multilayer perceptron (MLP), LSTM, and Gate Recurrent Unit (GRU)), the CNN model proposed in this study is promising.

As seen in the previous works, to predict COVID-19 cases, some works pretend that LSTM fits better than CNN, while others pretend the contrary. In this present work, we will use the CNN model in this context and compare our results with another submitted work investigating the efficiency of LSTM for the same objective on the same dataset.

### 3 MATERIALS AND METHODS

The approach consists in three principle tasks. The first one is a data pre-processing to prepare data to be analysed. The second, the third and the last one, respectively, are the model parametrization, training and testing. Those tasks are recursive according to the evaluation metrics until finding the best-fitted model. The retained model is then used for the prediction.

#### 3.1 Data Set

This For our study, we have decided to explore data from Our World in Data that uses three statistical resources on the COVID-19 pandemic published by the European Center for Disease Prevention and Control (ECDC). The dataset contains aggregated data from countries around the world that are available on GitHub. The variables represent data related to confirmed cases, deaths, and testing, as well as other variables of potential interest. The columns are: iso code, continent, location, date, total cases, new cases, total deaths, new deaths, total cases per million, new cases per million, total deaths per million, new deaths per million, total tests, new tests, new tests smoothed, total tests per thousand, new tests per thousand, new tests smoothed per thousand, tests units, stringency index, population, population density, median age, aged 65 older, aged 70 older, gdp per capital, extreme poverty, cardiovascular death rate, diabetes prevalence, female smokers, male smokers, handwashing facilities, hospital beds per thousand, life expectancy. Note that the analysed data belong to the period from March 02, 2020 to June 15, 2020.

#### 3.2 Pre-processing

The collected data were then processed by a Python project using the libraries Pandas, NumPy, SciPy and Matplotlib in a jupyter notebook.

We have first filtered data concerning Morocco, and then selected the features 'total\_cases', 'population', 'total\_deaths', and 'new\_deaths'. We have retained the last feature 'new\_cases' because it is the target of our predictions. The features concerning the total of deaths and cases are useful for the prediction because the contamination can arise from infected people that are always alive.

The third task was the feature scaling to normalize the range of independent variables or features of data so that each feature contributes approximately and proportionately in the ML algorithm. We have then applied the Min-Max scaler that transforms all values between the range 0 and 1. The fourth task of the pre-processing process is adapting our model to learn from the past time-steps in order to predict the positive COVID-19 cases for the future 7 time-steps, where we split the dataset into 80% of the training set and 20% of the testing set. The model uses two days lags as inputs to predict seven days as outputs.

### 3.3 Parameters

The choice of the activation function, the optimizer, and the number of layers were fixed at the beginning, while the number of filters in the CNN layers were updated recursively according to the evaluation steps to enhance the performance of the trained model. The best performance was obtained by 127 filters. The amount of days to return by the model depends on the number of outputs in last layer, in our paper, that is set to 7.

In this work, we have opted for the activation function ReLu and the Adam Optimizer belonging to the stochastic gradient descent SGD category.

In a neural network, the activation function takes in the output signal from the previous cell and converts it into some form that can be taken as input to the next cell. Similarly, in an artificial neural network (ANN), it transforms the summed weighted input from the node into the output for that input. The rectified linear activation function ReLU is a linear function that outputs the input directly if it is positive, otherwise, it will output zero. It is especially when using CNN, a widely used activation function and easy to compute which does not saturate, and prevents the Vanishing Gradient Problem.

The optimizer is a basic algorithm responsible for making neural networks converge, it controls the weights updating of the network, to shift towards the optimum of the cost function. On the basis of this, we have two basic types of gradient descent algorithms. The BGD, batch gradient descent and the SGD, stochastic gradient descent. The main distinction between the both is that the SGD will compute the cost of only one example in each step, when the BGD will determine the cost of all the training instances in the dataset.

The SGD formula is employed to adjust the weights in a neural network through the use backpropagation to calculate the gradient  $\nabla$  (Hansen, 2019):

$$\theta_{t+1} = \theta_t - \eta \cdot \overbrace{\nabla_{\theta} J(\theta; x, y)}^{\text{Backpropagation}} \quad (4)$$

- $\theta$  (theta) is the variable to be modified during the optimization of a model, it can be weights, biases and activations. Notice that a single weight is updated for the neural network here,
- $\eta$  is the learning rate (eta),
- $\nabla$  is the gradient (nabla),
- $J$  has formally been known as an objective function, but most often it is called a cost function or loss function.
- $J(\theta; x, y)$  basically implies that we insert the  $\theta$

parameter along with a training example and label.

Adaptive Moment Estimation (Adam) is the most frequently used optimizer and certainly the top performer on general average.

### 3.4 Evaluation Metrics

MSE (Mean Square Error). It is used as a default metric for evaluation of the performance of most regression algorithms (Kathuria, 2019). It computes the average of the squared errors, defined as the mean difference between the real values  $y$  and the estimated  $\hat{y}$  (Binieli, 2018). A large MSE means a large error.

RMSE (Root Mean Square Error) has been used as a standard statistical metric to measure model performance in meteorology, air quality, and climate research studies (Chai & Draxler, 2014). It corresponds to the square root of the mean of the squared difference between the observed  $y$  and the predicted values  $\hat{y}$  (Bratsas, et al., 2020).

Max Error is a metric which calculates the maximum residual error. It expresses the worst case error between the predicted value and the true value. Its interpretation is very simple.

R2 (R-Square) is a metric that varies between 0 and 1. It is also known as the Coefficient of Determination. Sometimes, the closer to 1 it is, the better the model is (Vedova, 2018).

MAE (Mean Absolute Error) is the averaged on the test instance for absolute differences between the forecast and the true observation where all individual errors were given the same weight.

### 3.5 1D-CNN Architecture

The 1D-CNN structure proposed in this study is illustrated in Figure 2 The design of the input layer uses two-time sequences that are based on factors influencing the number of new cases. The structure of the convolutional layer contains a single layer with 127 filters.

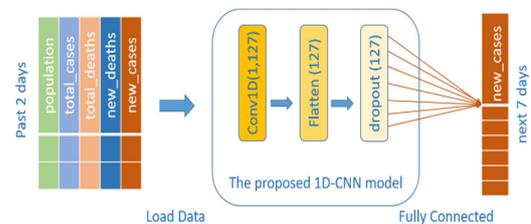


Figure 2: Structure of proposed CNN model.

The convolution process can be expressed in Equation (5). Let call  $f$  our input vector and  $g$  our

filter, and say that  $f$  has length  $n$ , and  $g$  has length  $m$ . The convolution  $f * g$  of  $f$  and  $g$  is defined as:

$$y_i = \sigma (w_1 x_1 + \dots + w_m x_m) \quad (5)$$

The idea behind the fully connected layer architecture can be formulated by equation (5). Let  $x \in \mathbb{R}^m$  represent the input to a fully connected layer. Let  $y_i \in \mathbb{R}$  be the  $i$ -th output from the fully connected layer. Then  $y_i \in \mathbb{R}$  is computed as follows:

$$x_i = (f * g)[i] = \sum_{j=1}^m g[j] * f[i - j + \frac{m}{2}] \quad (6)$$

Here,  $\sigma$  is a nonlinear function (in this case is ReLu), and the  $w_i$  are learnable weights (thanks to the optimizer) in the network. The result of the convolution layer can generate a 1D vector through flattened technique. Then the vectors are plugged into the fully connected (dense) layer to obtain a 1D vector output.

## 4 RESULTS AND DISCUSSION

In this section, we compare and discuss the results of the proposed CNN model and the best-fitted LSTM model developed by our research team. The two models were trained and tested on the same dataset. Table 1 shows the evaluations from the proposed CNN using the MSE, RMSE, MAE, R2, and MaxError metrics. As seen, the model converges well to a small error of the training loss with an MSE of 267.23, an RMSE of 16.35, an MAE of 11.74, and an R2 of 0.68. Actually, the model continues to well converge to a small error in the test stage with an MSE of 723.95, an RMSE of 26.91, an MAE of 20.06, and an R2 of 0.15.

Table 2 shows the experiment results of the best-fitted model among the LSTM ones that our research team has developed. It shows that the LSTM model converges slightly better than the proposed CNN model, in the test stage with an MSE of 658.65, an RMSE of 25.66, an MAE of 19.95, and an R2 of 0.23. Taking into consideration that the best-fitted LSTM model implements three LSTM layers, while the proposed CNN model implements only one CNN layer. It must be admitted that the proposed model is very promising.

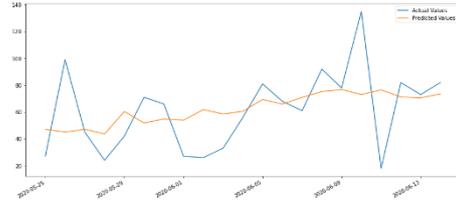


Figure 3: Curve of the CNN test.

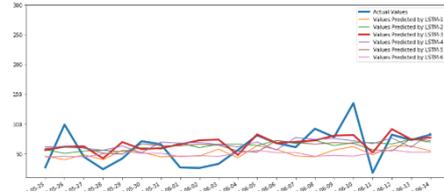


Figure 4: Test curves from research team LSTM models.

Examining these curves for the purpose of the comparison between real values and predicted values using the proposed CNN model (Figure 3) and the LSTM model (red one on Figure 4), we can notice that the two models are very close.

Table 1: Evaluation of the proposed CNN model.

	Model	MSE	RMSE	MAE	R2
Train	CNN	267.23	16.35	11.74	0.68
Test	CNN	723.95	26.91	20.06	0.15

Table 2: Test results of our research team best-fitted LSTM model.

	Model	MSE	RMSE	MAE	R2
Test	LSTM-3	658.65	25.66	19.95	0.23

## 5 CONCLUSIONS

In this work, we explore the efficiency of CNN model to predict new cases in Morocco using a dataset of time series data from March 02, 2020 to June 15, 2020. The experiments showed that a CNN model with a single CNN layer is almost as efficient as an LSTM model with three LSTM layers. It allows a good prediction accuracy with an RMSE of 26.91. Our findings and conclusions are demonstrated and enhanced by various illustrations we provide in this paper.

The results of the proposed CNN are very promising and show that CNN can be a potentially good alternative to LSTM model for similar problem. This indeed encourages exploring its efficiency with other settings such as more layers and more features.

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