



Article A Novel Fractional-Order Discrete SIR Model for Predicting COVID-19 Behavior

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Abstract: During the broadcast of Coronavirus across the globe, many mathematicians made several mathematical models. This was, of course, in order to understand the forecast and behavior of this epidemic's spread precisely. Nevertheless, due to the lack of much information about it, the application of many models has become difficult in reality and sometimes impossible, unlike the simple SIR model. In this work, a simple, novel fractional-order discrete model is proposed in order to study the behavior of the COVID-19 epidemic. Such a model has shown its ability to adapt to the periodic change in the number of infections. The existence and uniqueness of the solution for the proposed model are examined with the help of the Picard Lindelöf method. Some theoretical results are established in view of the connection between the stability of the fixed points of this model and the basic reproduction number. Several numerical simulations are performed to verify the gained results.

Keywords: SIR model; fractional-order discrete operators; stability; existence and uniqueness; Picard Lindelöf method; basic reproduction number

MSC: 26A33; 39A05; 92B05

1. Introduction

Across the centuries, the world has seen the dissemination of several diseases, which have caused extensive loss of lives. In the first quarter of 2020, the World Health Organization (WHO) declared the COVID-19 pandemic around the world [1]. From such time up to now, the total number of global deaths has surpassed the obstacle of 5 million deaths. The world was compelled to track many preventive strategies such as using masks, sanitizers and imposing quarantine.

In terms of a mathematician's view, it is still essential to establish a proper mathematical model that can describe the growth of the COVID-19 pandemic and its effects on communities, see [2–9]. As a matter of fact, the Susceptible-Infectious-Removed model (or simply SIR model) is deemed one of the most distinguished mathematical models, which can be typically employed to outline the dynamics of such pandemic and evaluate possible scenarios of infection. The SIR model can be extremely useful for assessing the effectiveness of numerous strategies. This model depends in its construction on a kind of ordinary differential equations that consider the total population with respect to its infection, the



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Copyright: © 2022 by the authors. Licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (https:// creativecommons.org/licenses/by/ 4.0/). persons who unfortunately die, and the direction of persons who recover after infection over a given time [10,11].

In mathematical modeling, the ordinary differential equations are commonly utilized to establish certain models of continuous-time mode, whereas the difference equations are, on the other hand, utilized to establish other models of discrete-time mode. Regarding the continuous-time models, we find that there are some pioneer works that have recently dealt with the COVID-19 disease in very practical manners. For instance, a novel version of timedependent SIR model was recently constructed in [12] to investigate the transmission rate of COVID-19 disease in various regions of India, where such model was employed to show the effect of the complete knockdown on the transmission rate of the COVID-19 infection in Indian districts with the help of ArcGIS 10.2 by preparing district-wise spatial distribution maps. In [13], another fractional-order SIR model was derived from the stochastic process of a continuous-time random walk that incorporates a time-since-infection dependence on both the infectivity and the recovery of the population. In [14], several infectious diseases, including COVID-19 in its SIR form, were analyzed with the application of mathematical modeling in the analysis of their spreading rates and treatments. In [15], certain random walk models for the daily time series that confirmed COVID-19 cases for different countries were introduced and discussed, where such model was derived from the evolutionary equation for a specified memory function that can provide some non-ergodic fields, which were evident in some available COVID-19 data. At the same time, the discrete-time models have been recently employed to assess and explore the infectious diseases instead of the continuous-time models, as pandemic data are accessible during the periods of discretetime [16]. In addition, the discrete-time models can demonstrate more rich dynamics than the continuous-time ones. In order to provide an overview of some kinds of discrete models, the reader may refer to [17-20]. In particular, a discrete SIR epidemic model with a constant vaccination strategy in terms of its dynamical behavior was examined in [21], whereas the authors in [22] investigated the bifurcations and stability of a proposed discrete-time model of a susceptible-infected-susceptible (SIS) mode with vaccination. In addition, another proposed discrete-time model with vaccination was examined in terms of its dynamical behavior by Xiang et al. in [23]. In the same regard, a new generalization version of the fractional-order SIR epidemic model for predicting the spread of the COVID-19 disease was presented in [24], where the time-domain model implementation was based on the fixed-step method using the nabla fractional-order difference Grünwald-Letnikov operator.

In mathematical analysis, fractional calculus is considered one of the most important subjects as a generalization of integer calculus [25–30]. It has been demonstrated that the fractional-order derivative is significantly more precise than the integer-order one [31]. This is because it plays a key role as a powerful tool for outlining the effect of memory on all kinds of materials and processing [32]. From the perspective of this view, a lot of researchers have, over the last several years, been concerned with discrete fractional calculus [33–37]. In fact, there are many applications of such field in reality, such as the applications connected with biology, physics, neural networks, etc.

In the past few years, many epidemiology SIR models have been established in view of fractional-order differential equations [38–40]. In particular, a fractional-order SIR model was investigated in terms of its stability in implementing a discretization scheme in [41]. Naik proposed another fractional-order SIR model in terms of its global dynamics in [42]. Several numerical simulations with their analysis were performed for a new fractional-order SIR model in [43]. A delayed incommensurate fractional-order SIR model was also examined in terms of its bifurcation control and stability analysis in [44]. In view of the above considerations, it should be emphasized that all the aforesaid studies were established based on a continuous-time mode. This actually motivated us to propose a new fractional-order discrete model to be considered in the epidemiology field. As far as we know, the study of fractional-order discrete-time COVID-19 models has not probably been explored till now. Thus, from this perspective, the main target of this work is to make a contribution to the epidemiology field by proposing a new fractional-order discrete-time

SIR model. In this regard, it is worth mentioning that due to the fact the proposed model is just a fractionalization of the discrete SIR-type system and due to the pandemic time series that can be generated from its dynamics, we believe that this model will, in general, have several potential applications and perspectives.

In fact, in order to formulate the target mathematical model, which will enable us later to predict the COVID-19 behavior according to the available information, we suppose that the society is divided into four different classes; susceptible class *S*, infected class *I*, recovered class *R* and death class *D*. In addition, we suppose that the transition between these classes takes place according to the scheme exhibited in Figure 1.

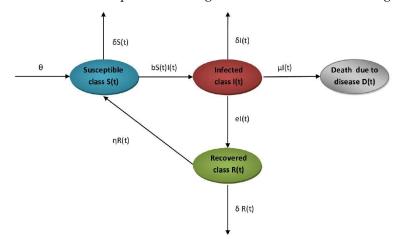


Figure 1. Disease transmission flow of the proposed model.

In light of this figure, the values of all parameters reported in the model at hand can be explained, as shown in Table 1.

Table 1. Description of the model's parameters.

Parameters	Description			
μ	Corona death rate			
δ	Natural death rate			
heta	The number of new births			
b	Infection rate			
е	Recovery rate			
η	The rate at which a recovering person is at risk of infection			

As a matter of fact, the disease transmission flow of the proposed fractional-order discrete SIR model, which is shown in its compartmental form in Figure 1, can be furthermore described by the following nonlinear system:

$$S(n+1) = S(n) + \theta + \eta R(n) - bS(n)I(n) - \delta S(n),$$

$$I(n+1) = I(n) + bS(t)I(n) - (\mu + \delta + e)I(n),$$

$$R(n+1) = R(n) + eI(n) - (\delta + \eta)R(n),$$

$$D(n+1) = D(n) + \mu I(n),$$

(1)

subject to the following initial conditions:

$$S(0), I(0), R(0), D(0) \ge 0.$$
 (2)

Since the first three equations in the proposed system (1) are not related to class D, and because this class can be studied alone from using the equation:

$$N(n) = S(n) + I(n) + R(n) + D(n),$$
(3)

then the last equation of system (1) might be ignored. This would actually facilitate our examination by focusing only on the first three equations of that system. At the same time, it should be noted that because the value of the parameter *b* is very small ($b = \frac{pk}{N}$, where *k* is the rate of contacts per unit of time so $0 \le k \le 1$, *p* is the probabilities of contagion so $0 \le p \le 1$, and *N* is the number of the total population in the tens of thousands or more), the two terms bS(n) and bI(n) are also very small. The fact that we did not mention the age groups and vaccination does not mean that they were neglected, but the high life expectancy in a society can be seen as one of the causes of the high death rate due to the virus. As for vaccination, we know that it does not provide 100 percent immunity, but it reduces the possibility of infection (which reduces *b*), reduces virus mortality (η), and increases the recovery rate (*e*).

The rest of this paper is organized as follows: Section 2 recalls some basic definitions and primary facts associated with discrete fractional calculus. In Section 3, we use the fixed point theory coupled with the Picard Lindelöf method for the purpose of showing the existence and uniqueness of the solution for the proposed fractional-order discrete-time COVID-19 model. In Section 4, we analyze the stability of the fixed points of the proposed model, and we find the basic reproduction number of an infectious disease. Section 5 shows several numerical simulations performed to verify the theoretical findings, followed by Section 6, which summarizes the conclusions of this work.

2. Preliminaries

During the past few decades, fractional calculus has demonstrated high efficiency in providing more practical outcomes than that of traditional calculus. This, actually, is due to its flexibility and ability to provide approximations of several phenomena better than before. However, in order to gain more depth in understanding the proposed fractional-order model in this work and its impact on reality, it is useful to outline some definitions and basic facts connected with discrete fractional calculus. From now on, we will assume that all functions are defined on the set $\mathbb{N}_a := \{a, a + 1, a + 2, \dots\}$, where $a \in \mathbb{R}$.

Definition 1 ([45]). The α^{th} -fractional sum of the function $f : \mathbb{N}_a \to \mathbb{R}$ is defined by:

$$\Delta_a^{-\alpha} f(t) = \frac{1}{\Gamma(\alpha)} \sum_{s=a}^{t-\alpha} (t-s-1)^{(\alpha-1)} f(s), \text{ for } t \in \mathbb{N}_{a+\alpha},$$
(4)

where $\alpha > 0$, $\Gamma(\cdot)$ is the Euler's gamma function and $t^{(\alpha)} = \frac{\Gamma(t+1)}{\Gamma(t+1-\alpha)}$.

Definition 2 ([45]). *The* α^{th} *-order Rimann–Liouville fractional difference operator of the function f defined on* \mathbb{N}_a *is outlined by:*

$$\Delta_{a}^{\alpha}f(t) := \Delta\Delta_{a}^{-(1-\alpha)}f(t) = \frac{1}{\Gamma(1-\alpha)}\Delta\sum_{s=a}^{t-(1-\alpha)}(t-s-1)^{(-\alpha)}f(s), \text{ for } t \in \mathbb{N}_{a+1-\alpha},$$
(5)

where $0 < \alpha < 1$.

Definition 3 ([45]). The α th-order Caputo fractional difference operator of the function f defined on \mathbb{N}_a is outlined by:

$${}^{C}\Delta_{a}^{\alpha}f(t) = \begin{cases} \Delta_{a}^{-(1-\alpha)}\Delta f(t) = \frac{1}{\Gamma(1-\alpha)} \sum_{s=a}^{t-(1-\alpha)} (t-s-1)^{(-\alpha)}\Delta f(s), & 0 < \alpha < 1 \\ \Delta f(t), & \alpha = 1, \end{cases}$$
(6)

where $0 < \alpha \leq 1$ and $t \in \mathbb{N}_{a+1-\alpha}$.

Proposition 1 ([45]). *Let f be a function defined on* \mathbb{N}_a *. Then:*

$$\Delta_{a+(1-\alpha)}^{-\alpha} {}^{C}\Delta_{a}^{\alpha} f(t) = f(t) - f(a), \quad \forall t \in \mathbb{N}_{a},$$
(7)

where $0 < \alpha \leq 1$.

Proposition 2 ([45]). *Let f be a function defined on* \mathbb{N}_a *. Then:*

$${}^{C}\Delta_{a}^{\alpha} f(t) = \Delta_{a}^{\alpha} f(t) - \frac{(t-a)^{(-\alpha)}}{\Gamma(1-\alpha)} f(a), \quad \forall t \in \mathbb{N}_{a-\alpha+1},$$
(8)

where $0 < \alpha < 1$.

Lemma 1 ([46]). Let $\alpha > 0$, $\alpha \notin \mathbb{N}$ and f be a function defined on \mathbb{N}_a . Then:

$$f(t) = f(a) + \frac{1}{\Gamma(\alpha)} \sum_{r=a+1-\alpha}^{t-\alpha} (t-r-1)^{(\alpha-1)} {}^{C} \Delta_{a}^{\alpha} f(r), \quad \forall t \in \mathbb{N}_{a+1}.$$
(9)

Remark 1. One might observe the following states:

- If ${}^{C}\Delta_{a}^{\alpha}f(t) \ge 0$, then the function f is nondecreasing for all $t \in \mathbb{N}_{a}$.
- If ${}^{C}\Delta_{a}^{\alpha}f(t) \leq 0$, then the function f is nonincreasing for all $t \in \mathbb{N}_{a}$.

Theorem 1 ([47]). Suppose h(t, y) and k(t, y) are two real-valued functions defined on $[0, +\infty) \times \mathbb{R}$. If the function k satisfies the Lipschitz condition in y with Lipschitz constant L_k such that $0 < L_k \le \alpha$, then $y_1(t)$ and $y_2(t)$ are, respectively, two unique solutions for the following initial value problems:

$$\begin{cases} \Delta_a^{\alpha} y(t) = h(t+\alpha-1, y(t+\alpha-1)), \ t \in \mathbb{N}_{a+1-\alpha} \\ \Delta_a^{\alpha-1} y(t)\big|_{t=a} = y_0, \end{cases}$$
(10)

and

$$\begin{cases} \Delta_a^{\alpha} y(t) = k(t+\alpha-1, y(t+\alpha-1)), \ t \in \mathbb{N}_{a+1-\alpha} \\ \Delta_a^{\alpha-1} y(t) \big|_{t=a} = y_0. \end{cases}$$
(11)

Herein, based on the above theorem, it can be noted the following observations:

- If h(t, y) < k(t, y), then $y_1(t) \le y_2(t)$ for $t \in \mathbb{N}_a$.
- If h(t,y) > k(t,y), then $y_1(t) > y_2(t)$ for $t \in \mathbb{N}_a$.

Remark 2. In view of Proposition 2, we can notice that Theorem 1 remains true if we replace Δ_a^{α} by ${}^{C}\Delta_a^{\alpha}$.

In accordance with the nonlinear model given in (3), we can now define the fractionalorder model, which will be taken into consideration in this work from now on. This model has the following form:

$$\begin{cases} {}^{C}\Delta_{0}^{\alpha}S(t+1-\alpha) = \theta + \eta R(t) - bS(t)I(t) - \delta S(t),\\ {}^{C}\Delta_{0}^{\alpha}I(t+1-\alpha) = bS(t)I(t) - (\mu + \delta + e)I(t),\\ {}^{C}\Delta_{0}^{\alpha}R(t+1-\alpha) = eI(t) - (\delta + \eta)R(t), \end{cases}$$
(12)

subject to the following initial conditions:

$$S(0), I(0), R(0), D(0) \ge 0,$$
 (13)

where $0 < \alpha < 1$ and $t \in \mathbb{N}$. Continuing to move forward in our examination, we recall below an important result that clarifies the stability of the solution of system (12).

Theorem 2 ([48,49]). *Let* $\alpha \in (0,1)$. *If*

$$\lambda \in \left\{ z \in \mathbb{C} : |z| < \left(2\cos\frac{|\arg z| - \pi}{2 - \alpha} \right)^{\alpha} and |\arg z| > \frac{\alpha \pi}{2} \right\},\tag{14}$$

for all the eigenvalues λ of *I*, then the fixed point of system (12) is asymptotically stable, where *I* is the Jacobite matrix of the second member of system (12) at the fixed point itself.

3. Existence and Uniqueness

In the following content, we intend to use the fixed point theory together with the Picard Lindelöf method to show the existence and uniqueness of solution for system (12). For this purpose, we may rewrite such system in the following classical form:

$$\begin{cases} {}^{C}\Delta_{0}^{\alpha}X(t) = F(t-1+\alpha, X(t-1+\alpha)), \\ X(0) = X_{0}, \end{cases}$$
(15)

(15) where $t \in \mathbb{N}_{1-\alpha}^{T_{\max}}$ in which $(T_{\max} - 1 + \alpha) \in \mathbb{N}$, $X(t) = (S(t), I(t), R(t), D(t))^T$ and the function F(t, X(t)) is defined as follows:

$$F_{1}(t,S) = \theta + \eta R(t) - bS(t)I(t) - \delta S(t), F_{2}(t,I) = bS(t)I(t) - (\mu + \delta + e)I(t), F_{3}(t,R) = eI(t) - (\delta + \eta)R(t).$$
(16)

In order to proceed with the existence and uniqueness investigation, we use the initial condition X(0) as well as Proposition 1. This would immediately transform system (12) into the following sum equations:

$$\begin{cases} S(t) - S(0) = & \Delta_{1-\alpha}^{-\alpha}(\theta + \eta R(t) - bS(t)I(t) - \delta S(t)), \\ I(t) - I(0) = & \Delta_{1-\alpha}^{-\alpha}(bS(t)I(t) - (\mu + \delta + e)I(t)), \\ R(t) - R(0) = & \Delta_{1-\alpha}^{-\alpha}(eI(t) - (\delta + \eta)R(t)), \end{cases}$$
(17)

for $t \in \mathbb{N}_{1-\alpha}^{T_{\max}}$. Using system (17) and Definition 1, we can obtain the state variable in terms of $F_i(t, X(t))$, where i = 1, 2, 3. In other words, we have:

$$\begin{cases} S(t) = S(0) + \frac{1}{\Gamma(\alpha)} \sum_{s=1-\alpha}^{t-\alpha} (t-s-1)^{(\alpha-1)} F_1(s-1+\alpha, S(s-1+\alpha)), \\ I(t) = I(0) + \frac{1}{\Gamma(\alpha)} \sum_{s=1-\alpha}^{t-\alpha} (t-s-1)^{(\alpha-1)} F_2(s-1+\alpha, I(s-1+\alpha)), \quad t \in \mathbb{N}_1^{T_{\max}}. \end{cases}$$
(18)
$$R(t) = R(0) + \frac{1}{\Gamma(\alpha)} \sum_{s=1-\alpha}^{t-\alpha} (t-s-1)^{(\alpha-1)} F_3(s-1+\alpha, R(s-1+\alpha)),$$

Thus, with the help of Picard iterations, we can obtain the following equations:

$$S_{n+1}(t) = S(0) + \frac{1}{\Gamma(\alpha)} \sum_{s=1-\alpha}^{t-\alpha} (t-s-1)^{(\alpha-1)} F_1(s-1+\alpha, S_n(s-1+\alpha)),$$

$$I_{n+1}(t) = I(0) + \frac{1}{\Gamma(\alpha)} \sum_{s=1-\alpha}^{t-\alpha} (t-s-1)^{(\alpha-1)} F_2(s-1+\alpha, I_n(s-1+\alpha)), \quad t \in \mathbb{N}_1^{T_{\max}}.$$

$$R_{n+1}(t) = R(0) + \frac{1}{\Gamma(\alpha)} \sum_{s=1-\alpha}^{t-\alpha} (t-s-1)^{(\alpha-1)} F_3(s-1+\alpha, R_n(s-1+\alpha)),$$
(19)

Consequently, based on system (17), together with its initial condition, we can gain the following sum equation:

$$X(t) = X(0) + \frac{1}{\Gamma(\alpha)} \sum_{s=1-\alpha}^{t-\alpha} (t-s-1)^{(\alpha-1)} F(s-1+\alpha, X(s-1+\alpha)),$$
(20)

where $t \in \mathbb{N}_1$. As a result of the previous groundwork, we can infer the following theoretical results that concern the existence and uniqueness of a solution for the system at hand.

Lemma 2. The function F(t, X(t)) defined in (16) satisfies the following Lipschitz condition:

$$\|F(t, X(t) - F(t, X^*(t))\| \le \beta \|(X(t) - X^*(t))\|,$$
(21)

where

$$\beta = \max \left\{ \begin{array}{c} \max \| bS(t) - (\mu + \delta + e) \|, \\ \max \| (\delta + bI(t)) \|, \| (\delta + \eta) \| \end{array} \right\}.$$
(22)

Proof. By taking S(t) and $S^*(t)$ as two coupled functions, we obtain:

$$\|F_1(t,S) - F_1(t,S^*)\| = \|(\delta + bI(t))(S - S^*)\|.$$
(23)

At the same time, by taking into account:

$$\beta_1 = \max \| (\delta + bI(t)) \|, \tag{24}$$

one can reach:

$$\|F_1(t,S) - F_1(t,S^*)\| \le \beta_1 \|(S - S^*)\|.$$
(25)

Continuing in a similar manner yields the following two inequalities:

$$\|F_2(t,I) - F_2(t,I^*)\| \leq \beta_2 \|(I-I^*)\|, \|F_3(t,R) - F_3(t,R^*)\| \leq \beta_3 \|(R-R^*)\|,$$
(26)

where

$$\beta_{2} = \max \|bS(t) - (\mu + \delta + e)\|, \beta_{3} = \|(\delta + \eta)\|.$$
(27)

From (24)–(26), we can confirm that the kernels F_1 , F_2 and F_3 satisfy the Lipschitz condition. Moreover, if $\beta_i < 1$, then the kernel F_i is a contraction for i = 1, 2, 3.

Theorem 3. Assume that condition (21) is satisfied, then there exists a unique solution of system (12) if:

$$\frac{(T_{\max} - 1 + \alpha)^{(\alpha)}}{\Gamma(\alpha + 1)}\beta < 1.$$
(28)

Proof. Actually, the solution of system (12) can be outlined as:

$$X(t) = P(X(t)), \tag{29}$$

where *P* is the Picard operator defined by:

$$P(X(t)) = X(0) + \frac{1}{\Gamma(\alpha)} \sum_{s=1-\alpha}^{t-\alpha} (t-s-1)^{(\alpha-1)} F(s-1+\alpha, X(s-1+\alpha)).$$
(30)

Based on the previous arguments, we can have:

$$\begin{split} |P(X_{1}(t) - P(X_{2}(t))|| &= \left\| \frac{1}{\Gamma(\alpha)} \sum_{s=1-\alpha}^{t-\alpha} (t-s-1)^{(\alpha-1)} (F(s-1+\alpha,X_{1}(s-1+\alpha)) \\ &-F(s-1+\alpha,X_{2}(s-1+\alpha))) \right\|, \\ &\leq \frac{1}{\Gamma(\alpha)} \sum_{s=1-\alpha}^{t-\alpha} (t-s-1)^{(\alpha-1)} \| (F(s-1+\alpha,X_{1}(s-1+\alpha)) \\ &-F(s-1+\alpha,X_{2}(s-1+\alpha))) \|, \\ &\leq \frac{1}{\Gamma(\alpha)} \left(\sum_{s=1-\alpha}^{t-\alpha} (t-s-1)^{(\alpha-1)} \right) \max_{s\in\mathbb{N}_{1-\alpha}^{t-\alpha}} \| (F(s-1+\alpha,X_{1}(s-1+\alpha)) \\ &-F(s-1+\alpha,X_{2}(s-1+\alpha))) \|, \\ &\leq \frac{(t-1+\alpha)^{(\alpha)}}{\Gamma(\alpha+1)} \beta \| (X_{1}(t) - X_{2}(t)) \|. \end{split}$$

Since $\frac{(T_{\max}-1+\alpha)^{(\alpha)}}{\Gamma(\alpha+1)}\beta < 1$, then the operator *P* is a contraction, where $t \leq T_{\max}$. Hence, system (12) has a unique solution. \Box

In this regard and based on the aforesaid discussion, we can deduce, for the population size N = S + I + R, the following assertion:

$${}^{C}\Delta_{a}^{\alpha}N(t+1-\alpha) = \theta - \delta N(t) - (\mu + \gamma)I(t) \le \theta - \delta N(t).$$
(31)

This, consequently, yields:

$${}^{C}\Delta_{a}^{\alpha}N(t+1-\alpha) \leq {}^{C}\Delta_{a}^{\alpha}Y(t+1-\alpha),$$
(32)

where

$$\begin{cases} {}^{C}\Delta_{a}^{\alpha}Y(t+1-\alpha) = \theta - \delta Y(t),\\ Y(a) = N(a). \end{cases}$$
(33)

Thus, with the help of using Theorem 1 and Remark 2, it follows that:

$$N(t) \le Y(t),\tag{34}$$

that is because $N(a) \leq \frac{\theta}{\delta}$. Now, if we assume $t^* \in \mathbb{N}_a$ is the first point, where $Y(t^*) > \frac{\theta}{\delta}$, then we have:

$$\begin{split} Y(t^*) &= Y(a) + \frac{1}{\Gamma(\alpha)} \sum_{\substack{r=a+1-\alpha}}^{t^*-\alpha} (t^*-r-1)^{(\alpha-1)-C} \Delta_a^{\alpha} Y(r) \\ &= Y(a) + \frac{1}{\Gamma(\alpha)} \sum_{\substack{r=a+1-\alpha}}^{t^*-1-\alpha} (t^*-r-1)^{(\alpha-1)-C} \Delta_a^{\alpha} Y(r) + \frac{(\alpha-1)^{(\alpha-1)}}{\Gamma(\alpha)} C \Delta_a^{\alpha} Y(t^*-\alpha) \\ &= Y(a) + \frac{1}{\Gamma(\alpha)} \sum_{\substack{r=a+1-\alpha}}^{t^*-1-\alpha} (t^*-r-1)^{(\alpha-1)-C} \Delta_a^{\alpha} Y(r) + \frac{(\alpha-1)^{(\alpha-1)}}{\Gamma(\alpha)} C \Delta_a^{\alpha} Y(t^*-\alpha) \\ &= Y(a) + \frac{1}{\Gamma(\alpha)} \sum_{\substack{r=a+1-\alpha}}^{t^*-1-\alpha} \frac{(t^*-r-1)}{(t^*-r-\alpha)} (t^*-r-2)^{(\alpha-1)-C} \Delta_a^{\alpha} Y(r) + \frac{(\alpha-1)^{(\alpha-1)}}{\Gamma(\alpha)} C \Delta_a^{\alpha} Y(t^*-\alpha) \\ &= Y(t^*-1) + \frac{1}{\Gamma(\alpha)} \sum_{\substack{r=a+1-\alpha}}^{t^*-1-\alpha} \frac{(\alpha-1)}{(t^*-r-\alpha)} (t^*-r-2)^{(\alpha-1)-C} \Delta_a^{\alpha} Y(r) + \frac{(\alpha-1)^{(\alpha-1)}}{\Gamma(\alpha)} C \Delta_a^{\alpha} Y(t^*-\alpha) \\ &\leq Y(t^*-1) + \theta - \delta Y(t^*-1) \\ &\leq \frac{\theta}{\delta}, \quad \text{if } \delta < 1. \end{split}$$

Thus, we have a contradiction. Therefore, from this result, we can also prove that if $\delta < 1$, then we can have:

$$Y(t) \ge 0. \tag{35}$$

This, actually, asserts that the solution of the system belongs to the region:

$$\left\{ X \in \mathbb{R}^3, \ 0 \le x_1 + x_2 + x_3 \le \frac{\theta}{\delta} \right\}.$$
(36)

It is worth mentioning that the solution of System (3) is positive when b is small enough. However, it can not be shown that the solution of system (12) is positive. Actually, the solution of this system sometimes has negative values, but it can then increase to have positive values once again. For example, if we let t^* be the first point at which *S* becomes negative, we obtain:

$${}^{C}\Delta_{0}^{\alpha}S(t^{*}+1-\alpha) = \theta - \delta S(t^{*}) - bS(t^{*})I(t^{*}) + \eta R(t^{*}) > 0.$$
(37)

Thus, in accordance with Remark 1, one can notice that the class *S* will then be increased immediately. This assertion can be, in a similar manner, proven for the two classes *I* and *R*. Although this matter is considered a paradox because the values of the system should not be negative, what has been observed is that this system can give more realistic results over time. However, this inference will be discussed later on.

4. Fixed Points and Stability Analysis

In this section, we will be concerned with analyzing the stability of the disease-free fixed point by finding sufficient conditions connected with the parameters of system (12) to ensure the stability of this point. To this aim, we will first find those fixed points by equating the right-hand side of system (12) with zero. In other words, we have:

$$\begin{cases} \theta + \eta R^* - bS^* I^* - \delta S^* = 0, \\ bS^* I^* - (\mu + \delta + e)I^* = 0, \\ eI^* - (\delta + \eta)R^* = 0. \end{cases}$$
(38)

It can be then seen that the above system has two fixed points at most. The first one is called the disease-free fixed point (or simply DFF point), which can be obtained by assuming $I^* = 0$, i.e.,

$$E_0 = \left(\frac{\theta}{\delta}, 0, 0\right). \tag{39}$$

On the other hand, if we let $I^* \neq 0$, then we obtain:

Ι

$$R^* = \frac{e}{(\delta + \eta)} I^*$$

$$S^* = \frac{(\mu + \delta + e)}{b},$$

$$(40)$$

$$S^* = \left(\frac{\delta(\mu + \delta + e) - b\theta}{b}\right) \left(\frac{\delta(\delta + \eta)}{\eta e - (\delta + \eta)(\mu + \delta + e)}\right).$$

This, actually, yields the second fixed point of the considered system, which is called the pandemic fixed point E^* . In other words, this point can be yielded only when $I^* > 0$, and it has the form $E^* = (S^*, I^*, R^*)$, where S^* , I^* and R^* are defined above.

In the same regard, the so-called basic reproduction number R_0 can be defined as the number of infections caused by the first disease case. This 'infections' number typically appears in an appointed population where everyone is assumed to be susceptible to infection [40]. The importance of R_0 lies in knowing the rapidity of the spread of the emerging disease among the inhabitants and the proportion of the population to be immunized [40]. To be precise, the population spread of the epidemic will occur when $R_0 > 1$, where it is difficult to control. The method to enumerate the basic reproduction number can be performed by finding the spectral radius of the next generation matrix Y (i.e., $R_0 = \rho(Y)$). The matrix Y is a multiplication of F by V^{-1} , where:

$$F = \begin{bmatrix} \frac{\partial F_i(E_0)}{\partial t_j} \end{bmatrix} \text{ and } V = \begin{bmatrix} \frac{\partial V_i(E_0)}{\partial t_j} \end{bmatrix}, \tag{41}$$

where F_i refers to the stream of freshly infected cases into compartment t_j , and V_i refers to the entering/leaving streams connected with t_j , for $i, j = 1, 2, 3, \dots, m$ such that m is the total of compartments demonstrated in the model. Based on the aforesaid argument, one might calculate R_0 for the fractional-order model (12) by obtaining the two primary

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matrices *F* and *V*. To this aim, we can note that the infected compartment *I* can yield the following assertion:

$$^{C}\Delta_{a}^{\alpha}I(t+1-\alpha) = bS(t)I(t) - (\mu+\delta+e)I(t).$$
(42)

This consequently implies the following Jacobian matrix:

$$J = \left(\frac{b\theta}{\delta}\right) - (\mu + \delta + e). \tag{43}$$

Accordingly, the above matrix can be decomposed in terms of the two matrices *F* and *V* so that J = F - V, where:

$$F = \left(\frac{b\theta}{\delta}\right),\tag{44}$$

and

$$V = (\mu + \delta + e). \tag{45}$$

Therefore, the basic reproduction number R_0 can be then calculated to be given as:

$$R_0 = \rho(Y) = \rho(FV^{-1}) = \frac{b\theta}{\delta(\mu + \delta + e)}.$$
(46)

Remark 3. It should be noted that:

$$I^* = \frac{(\delta + \eta)\theta}{(\mu + \delta + e)(\delta + \eta) - \eta e} \left(1 - \frac{1}{R_0}\right),\tag{47}$$

which asserts that the pandemic fixed point E^* will hold only when $R_0 > 1$.

In what follows, we will analyze the stability of the DFF point. This will be implemented by establishing some sufficient conditions related to the parameters of system (12) to ensure the stability of this point. In order to achieve this objective, we introduce the next theoretical result.

Theorem 4. In case of $R_0 < 1$, the DFE point (E_0) of system (12) is locally asymptotically stable if

$$\max\{(\delta + \mu + e)(1 - R_0), (\delta + \eta)\} < 2^{\alpha}.$$
(48)

Proof. The Jacobian matrix of *F* at E_0 can be given as:

$$J(E_0) = \begin{pmatrix} -\delta & -\frac{b\theta}{\delta} & \eta \\ 0 & \frac{b\theta}{\delta} - (\mu + \delta + e) & 0 \\ 0 & e & -(\delta + \eta) \end{pmatrix}.$$
 (49)

Consequently, the characteristic polynomial will be as:

$$\det(\lambda Id - J(E_0)) = (\lambda + \delta)(\lambda + \delta + \eta) \left(\lambda + \frac{\delta e - b\theta + \delta^2 + \mu\delta}{\delta}\right),$$

 \Leftrightarrow

$$\det(\lambda Id - J(E_0)) = (\lambda + \delta)(\lambda + (\delta + \eta))(\lambda - (\delta + \mu + e)(R_0 - 1))$$

This implies:

$$\det(\lambda Id - J(E_0)) = 0,$$

$$\lambda_1 = -\delta < 0 \tag{50}$$

$$\lambda_2 = -(\delta + \eta) < 0 \tag{51}$$

 \Leftrightarrow

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and

$$\lambda_3 = (\delta + \mu + e)(R_0 - 1) < 0.$$
(52)

Thus, when $R_0 < 1$, we have:

$$\begin{aligned} -2^{\alpha} < \lambda_1 < 0\\ -2^{\alpha} < \lambda_2 < 0. \end{aligned} \tag{53}$$

Hence, according to Theorem 2, the DFE point is locally asymptotically stable if $R_0 < 1$. \Box

5. Application to Predict the Behavior of COVID-19 in Germany

In this section, we will perform several numerical simulations to verify the results inferred in the previous sections. For this purpose, we will apply our study to predict the behavior of the virus in Germany. We will take all the statistics of a million people, which means we will take the initial population size N(0) as 1,000,000. According to [50], we can easily find: the number of new births per day for a million people $\theta = 26.3308$ and the death rate $\delta = 3.15 \times 10^{-5}$. We can also obtain the stats described in Table 2 from the site [51].

 Table 2. Real data: The number of active infections in Germany in the period 26 April to 23 May 2020 [51].

-	26-Apr	27-Apr	28-Apr	29-Apr	30-Apr	1-May	2-May
	30,791	29,637	28,642	28,126	27,845	27,375	26,262
-	3-May	4-May	5-May	6-May	7-May	8-May	9-May
-	25,884	25,693	24,826	24,234	23,968	23,565	22,531
_	10-May	11-May	12-May	13-May	14-May	15-May	16-May
_	21,817	21,253	20,707	20,664	20,557	20,250	19,914
	17-May	18-May	19-May	20-May	21-May	22-May	23-May
_	19,200	18,254	17,411	16,687	16,435	16,151	15,092

Using Table 2 and [51], we find the contact rate $b = 5.1160 \times 10^{-7}$, the recovery rate e = 0.0789, the rate at which a recovering person is at risk of infection $\eta = 0.87$, and the corona death rate $\mu = 0.1$. We also find that the initial number of the susceptible people is S(0) = 686,400, while the initial number of the exposed people is I(0) = 33,966, and finally, the initial number of the infected people is R(0) = 279,630. In order to apply Theorem 4, we first calculate the basic reproduction number:

$$R_0 = \frac{b\theta}{\delta(\mu + \delta + e)} = 0.73013 < 1, \tag{54}$$

and

$$\max\{(\delta + \mu + e)(1 - R_0), (\delta + \eta)\} < 2^{\alpha} = 0.87003 < 2^{\alpha}.$$
(55)

Note that the conditions of Theorem 4 are validated, then the DFE point is locally asymptotically stable. Anyhow, based on these, we plot Figures 2 and 3, which represents a numerical simulation, confirming the stability of the system (12) in this case.

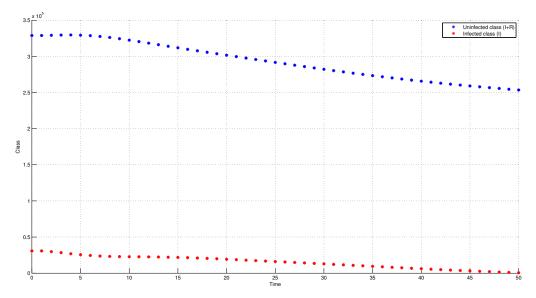


Figure 2. Numerical simulation of system (12).

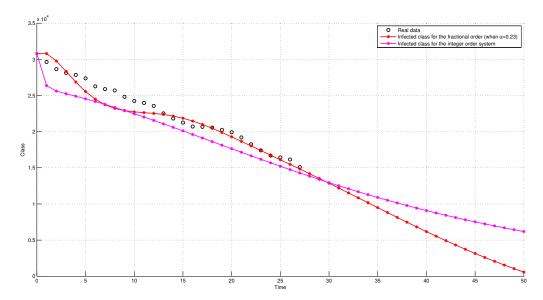


Figure 3. Numerical simulation of infected class with integer and commensurate fractional-order and comparison with the real data.

In order to obtain a better modeling use the incommensurate order system

$$\begin{cases} {}^{C}\Delta_{0}^{\alpha_{1}}S(t+1-\alpha) = \theta + \eta R(t) - bS(t)I(t) - \delta S(t), \\ {}^{C}\Delta_{0}^{\alpha_{2}}I(t+1-\alpha) = bS(t)I(t) - (\mu + \delta + e)I(t), \\ {}^{C}\Delta_{0}^{\alpha_{3}}R(t+1-\alpha) = eI(t) - (\delta + \eta)R(t), \end{cases}$$
(56)

where $0 \le \alpha_1, \alpha_2, \alpha_3 \le 1$. In this case, the existence of a direct and simple condition that deals with the stability of the system (56) is difficult and may be impossible in comparison with establishing a simple condition that deals with the stability of the commensurate system (12). However, it can indeed study the stability of the fixed points for system (56) by knowing the values of $\alpha_i, 1 \le i \le 3$. In such a case, we need to report the next corollary.

Corollary 1 ([48]). Suppose that $0 < \alpha_i < 1$, for $1 \le i \le 3$, and M is the least common multiple of u_i and v_i so that $\alpha_i = \frac{v_i}{u_i}$ with $gcd(u_i, v_i) = 1$ and $1 \le i \le 3$, where $gcd(\cdot, \cdot)$ is denoted to the greatest common divisor. If at least one root of the following equation:

$$\det\left(diag(\lambda^{M\alpha_1},\lambda^{M\alpha_2},\lambda^{M\alpha_3})-\left(1-\lambda^M\right)J\right)=0,$$
(57)

lies inside the set

$$\mathbb{C}\setminus K^{\frac{1}{M}}$$
, (58)

then the trivial solution of system (56) corresponding to (S(0), E(0), I(0)) is locally asymptotically stable, where

$$K^{\frac{1}{M}} = \left\{ z \in \mathbb{C} : |z| \le (2\cos(M|\arg z|))^{\frac{1}{M}} \text{ and } |\arg z| \le \frac{\pi}{2M} \right\}.$$
(59)

Overall we will consider $\alpha_1 = 0.223$, $\alpha_2 = 0.2$, $\alpha_3 = 0.232$. In this case, we obtain a more accurate approximation than the commensurate order case. Figure 4 shows a numerical simulation of this case, and Figure 5 represents the comparison of this case with the real data, while Figure 6 represents the comparison between the integer order case, the commensurate order case, the incommensurate order case, and the real data. It can be seen that each time, the system is more accurate.

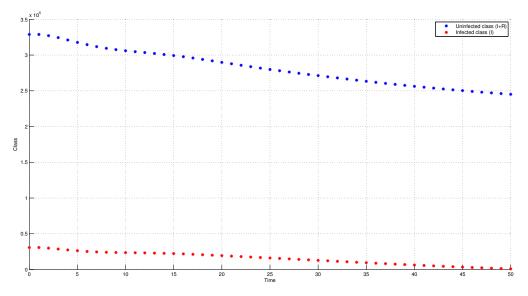


Figure 4. Numerical simulation of system (56) with ($\alpha_1 = 0.223, \alpha_2 = 0.2, \alpha_3 = 0.232$ (56).

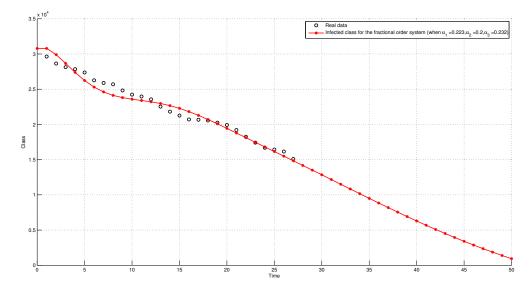
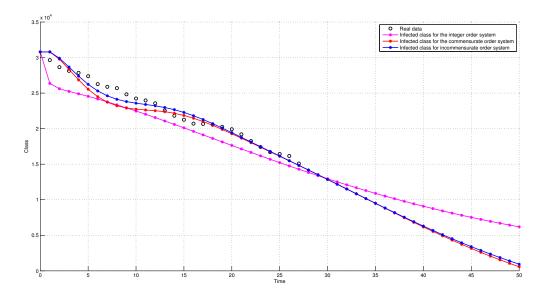


Figure 5. The comparison between the results of (56) and the real data.



It can be seen that system (56) is more accurate than system (12), and this reduces the error rate in prediction.

Figure 6. The comparison between the integer order, the commensurate order, and the incommensurate order systems with the real data.

6. Conclusions

In this work, a novel fractional-order discrete model has been proposed with the aim of studying the behavior of the COVID-19 pandemic. With the help of using the Picard Lindelöf method, the existence and uniqueness of the solution for the proposed model have been investigated. In light of the relationship between the stability of the fixed points of this model and the basic reproduction number, some useful theoretical results have been established. To verify these results, several numerical simulations have been performed via MATLAB scripts. In general, we can assert that the presented model in this work can be applied to study the COVID-19 pandemic in many regions around the world, even if this model is formulated with the fractional-order backward operator in its commensurate or incommensurate cases.

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