Optimality of Solution with Numerical Investigation for Coronavirus Epidemic Model

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Abstract: The novel coronavirus disease, coined as COVID-19, is a murderous and infectious disease initiated from Wuhan, China. This killer disease has taken a large number of lives around the world and its dynamics could not be controlled so far. In this article, the spatio-temporal compartmental epidemic model of the novel disease with advection and diffusion process is projected and analyzed. To counteract these types of diseases or restrict their spread, mankind depends upon mathematical modeling and medicine to reduce, alleviate, and anticipate the behavior of disease dynamics. The existence and uniqueness of the solution for the proposed system are investigated. Also, the solution to the considered system is made possible in a well-known functions space. For this purpose, a Banach space of function is chosen and the solutions are optimized in the closed and convex subset of the space. The essential explicit estimates for the solutions are investigated for the associated auxiliary data. The numerical solution and its analysis are the crux of this study. Moreover, the consistency, stability, and positivity are the indispensable and core properties of the compartmental models that a numerical design must possess. To this end, a nonstandard finite difference numerical scheme is developed to find the numerical solutions which preserve the structural properties of the continuous system. The M-matrix theory is applied to prove the positivity of the design. The results for the consistency and stability of the design are also presented in this study. The plausibility of the projected scheme is indicated by an appropriate example. Computer simulations are also exhibited to conclude the results.

Keywords: Spatio-temporal; advection; diffusion; explicit estimates; auxiliary data; M-matrix

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1 Introduction

Transmission of different infectious diseases occur in various parts of the world. To understand the dynamics of these, mathematicians have developed and analysed the mathematical models by taking into account their different aspects [1–7]. These models have a vital role to mitigate and to design the control strategy for the diseases. COVID-19 is one of the infectious diseases which creates severe acute respiratory syndrome. In December 2019, the first case of novel coronavirus was diagnosed in Wuhan, China and now, the whole world is in its deadly grip [8]. World Health Organization (WHO) declared this disease, a pandemic in the first half of 2020 [9]. According to the medical experts, this type of virus is extremely dangerous for old people having problems related to lungs, immune system, diabetes and blood pressure [10]. After the spread of this disease, the confirmed cases increased rapidly because of the non-availability of the vaccine. There may be more reasons for its spread, for instance, lack of awareness, lack of facilities to take the precautionary measures and many more. Due to this situation, the number of deaths are increasing day by day. The health workers including doctors, paramedical and other staff of the local health centers are in a high danger zone. Because, they face the infected people on the front lines. So, the identification of the infected cases is necessary at the initial stage so that the disease could not be transmitted to other people. Since the beginning of 2020, a large number of confirmed cases exposed in the other cities of China. Now, it has diffused all over the world. In the middle of April 2020, the virus affected almost all the countries on the globe and 1,929,995 confirmed cases and 119,789 deaths were recorded [11]. There are different thoughts in the world about the dispersion of COVID-19. According to some people, it is transmitted due to bats, and some associate it with the seafood market [12]. Another strong reason for the fast-spread of this virus is the traveling of infected persons, globally [13,14]. It is considered that COVID-19 is transmitted from animal to human, as most of the initial infected cases in Wuhan suspected that it is due to the seafood and animal market in the city [15], later, it was noticed that, the virus is transmitting from human to human also [16]. Now, in the current situation, this pandemic has become a great threat for the whole world. It has damaged seriously, the life of humans. Since harmful impacts can be seen in the economy, health, education, employment and social life. In order to provide security, every country is taking some serious steps depending on its available resources. The mathematicians are also observing the dynamics of this disease and trying to describe its behavior by developing the mathematical models. These models are helpful for the policymakers and scientists to understand the disease dynamics. They can make a future prediction of the disease communication and some precautionary measures may be adopted to save the lives of the population. The different aspects of COVID-19 has also attracted the other researchers and they have been studying the disease since a few months [17,18]. After developing a suitable mathematical model of this infection, it is a challenge for the researchers to find the solutions to these models. Usually, these models have a number of parameters (defining the rates with which various compartments are affected) so, in general, it becomes hard to find their analytical solutions. In this situation, the numerical solutions to these models are investigated. Numerical solutions and simulations help us to understand the behavior of the disease, especially when advection and diffusion terms are included in the model. In this article, a Spatio-temporal epidemic model of COVID-19 with advection and diffusion phenomena is analyzed. In the first section, the optimal existence of the solution is discussed and explicit estimates of the solution for the admissible auxiliary data are formulated. In Section 2, the non-standard finite difference (NSFD) method is developed for the proposed model and numerical solutions are found. In Section 3, the important physical properties of our numerical scheme are discussed. The positivity
of the proposed scheme is also proved by applying the M-matrix theory. In Section 4, numerical simulations are presented. Concluding remarks are given in the last Section 5. In this article, a Spatio-temporal epidemic model regarding COVID 19 with advection and diffusion is analyzed. The model is described as:

\[
\begin{align*}
S_t(x,t) + \alpha_1 S_x(x,t) &= a - cIS(1 + \gamma I) - \mu S + aR + d_1 S_{xx}(x,t) \\
I_t(x,t) + \alpha_2 I_x(x,t) &= cIS(1 + \gamma I) - (\beta + \mu + \delta - b)I + d_2 I_{xx}(x,t) \\
R_t(x,t) + \alpha_3 R_x(x,t) &= \beta I - (\alpha + \mu) R + d_3 R_{xx}(x,t)
\end{align*}
\]

(1)

where, \(S(t)\) represents the susceptible individuals, \(I(t)\) is identified as infected population, \(R(t)\) is recovered population, \(a\) represents the recruitment rate, \(\mu\) is the natural death rate, \(\delta\) (death rate due to corona virus), \(b\) (immigration rate of infected population), \(\alpha\) (recovery rate from infection), \(\beta\) (recovery rate from corona virus), \(c\) (infection rate) and \(\gamma\) describes the rate of recovered individuals that lose immunity. Also, \(\alpha_1, \alpha_2\) and \(\alpha_3\) represent the advection coefficients while \(d_1, d_2\) and \(d_3\) stand for diffusion coefficients.

The initial conditions for the system (1) are \(S(x,0) = S_0(x) > 0, I(x,0) = I_0(x) > 0\) and \(R(x,0) = R_0(x) > 0\).

The corona virus free equilibrium for the model (1) is \(E_0 = (\frac{\alpha}{\mu}, 0, 0)\).

And, the corona existence equilibrium is

\[
E_e = (S^*, I^*, R^*) = \left( \left( \frac{\beta}{\mu (\alpha + \mu)} - \frac{1}{\mu} (\beta + \mu + \delta - b) \right) I^* \right.
\]

\[
\left. - \frac{a}{\mu}. \frac{- (C\gamma + B) + \sqrt{(C\gamma - B)^2 + 4BA\gamma}}{2 (C - A)} \cdot \frac{\beta I^*}{(\alpha + \mu)} \right).
\]

where \(A = \mu (\alpha + \mu)(\beta + \mu + \delta - b), B = (\alpha c\beta - c (\alpha + \mu)(\beta + \mu + \delta - b))\) and \(C = -ac(\alpha + \mu)\).

The basic reproduction number denoted by \(R_0\), is given as \(R_0 = \frac{ac}{\mu (\beta + \mu + \delta - b)}\).

To examine the epidemic model (system of differential equations) incorporating the advection and diffusion terms, the understanding of the differential equations in epidemic model of the underlying system is necessary. These governing equations make the model easy for studying. To solve these equations for the underlying epidemic model numerically, it is important to investigate the existence, uniqueness, and convergence of the solution to the fixed point or equilibrium point [19,20]. Existence of the solution for these types of models is mandatory before finding the numerical solution. When the solution of the system is confirmed, then the next task is to find the desirable solutions that can be approximated. Lastly, a most feasible solution is picked out, which is in accordance with the constraints of the system. In general, the solutions to the differential equations lie in the spaces of functions which are known as Banach spaces. Generally, the solutions (functions) to the differential equations are not globally bounded so, it is always better to consider the subset of a function space. The explicit estimates can also be formulated. In 2005, Tutschke introduced the notion of explicit estimates for the general operator equations [21]. Initially, the problems are changed in to the form of fixed-point operators and their solutions are ensured in the subsets of the function spaces that lead to the restriction on the diameter of the subsets for given boundary values or intervals. Similarly, a chosen diameter can also lead to the restriction to the admissible auxiliary data. The subset (closed ball) for which the restriction

\[
\begin{align*}
S_t(x,t) + \alpha_1 S_x(x,t) &= a - cIS(1 + \gamma I) - \mu S + aR + d_1 S_{xx}(x,t) \\
I_t(x,t) + \alpha_2 I_x(x,t) &= cIS(1 + \gamma I) - (\beta + \mu + \delta - b)I + d_2 I_{xx}(x,t) \\
R_t(x,t) + \alpha_3 R_x(x,t) &= \beta I - (\alpha + \mu) R + d_3 R_{xx}(x,t)
\end{align*}
\]
is imposed, is called optimal ball. In the best of our knowledge, the explicit estimates with the optimal balls for any epidemic model have never been discussed.

2 Optimal Existence

In this section, optimal conditions are investigated in the frame work of Banach space, for some related quantities used in the procedure of developing the results, concerning the existence of optimal solutions of the proposed COVID-19 model (1). The system (1) can be reshaped in more general form

\[
\begin{align*}
S_t(x,t) &= G_1(S, I, R, S_x, S_{xx}) \\
I_t(x,t) &= G_2(S, I, R, I_x, I_{xx}) \\
R_t(x,t) &= G_3(S, I, R, R_x, R_{xx})
\end{align*}
\]

(2)

In this form \( G_1, G_2 \) and \( G_3 \) are real valued functions that may be nonlinear, such that the nonlinearity of \( G_1, G_2 \) and \( G_3 \) may depend not only on \( S, I \) and \( R \) but also on \( S_x, S_{xx}, I_x, I_{xx}, R_x \) and \( R_{xx} \). The solutions of above system (2) can be written in the following equivalent form:

\[
\begin{align*}
S(x,t) &= S_0 + \int_0^t G_1(S, I_0 + \int_0^\omega G_2(S, I, I_x, I_{xx}) (x, u) du, R_0 \\
&\quad + \int_0^\omega G_3(S, I, R_x, R_{xx}) (x, v) dv, S_x, S_{xx}) (x, \omega) d\omega.
\end{align*}
\]

According to the operator theory, the above integral equation which is considered for the solution of (2), can be written in operator form, as

\[
\mathcal{H}(S(x,t)) = S(x,t), S(x,t) \in C_{x,t}^{2,1}[a,b].
\]

Now, consider the Banach space \( X = C[0, \lambda] \), for any \( \lambda > 0 \). For the existence of the solution of system (1), Shauder fixed point theorem will be used here. For this, it is necessary to prove the fixed-point operator \( \mathcal{H} \) to be a self mapping on \( \overline{B(r, \Theta)} \), that is \( \mathcal{H} : \overline{B(r, \Theta)} \rightarrow \overline{B(r, \Theta)} \) defined by

\[
\mathcal{H}(S(x,t)) = S_0 + \int_0^t G_1(S, I_0 + \int_0^\omega G_2(S, I, I_x, I_{xx}) (x, u) du, R_0 \\
&\quad + \int_0^\omega G_3(S, I, R_x, R_{xx}) (x, v) dv, S_x, S_{xx}) (x, \omega) d\omega.
\]


Suppose also, that $\mathcal{H}$ is bounded by $\phi(r)$, that is $\|\mathcal{H}(x,t)\| \leq \phi(r)$, where $\phi(r)$ is a continuous real valued function. Optimization for the proposed model will be done in this space. For this purpose, a closed and convex subset of the Banach space will be chosen which is called a closed ball in $X$. The consideration of this closed ball yields two cases:

**Case 1.** Assume that $\overline{B}(r,\Theta) = \{ S \in X : \| S - \Theta \| \leq r, r > 0 \}$ be a closed ball with center at $\Theta$, zero of $X$, and radius $r$. Consider,

$$\|\mathcal{H}(x,t) - \Theta\| = \left\| S_0 + \int_0^t G_1(x,\omega)d\omega \right\|,$$

where,

$$G_1 \equiv G_1 \left( S, I_0 + \int_0^\omega G_2 (S, I, R, I_x, I_{xx}) (x,u)du, R_0 + \int_0^\omega G_3 (S, I, R, R_x, R_{xx}) (x,v)dv, S_x, S_{xx} \right)$$

$$\Rightarrow \|\mathcal{H}(x,t) - \Theta\| \leq |S_0| + \phi(r)t \leq |S_0| + \phi(r)\lambda$$

For $\mathcal{H}$ to be a self-mapping it should satisfy

$$|S_0| + \phi(r)\lambda \leq r. \quad (3)$$

The optimized value of $r$ can be obtained by fixing $S_0$ and $\lambda$, the length of interval of continuity, which is calculated from

$$r \geq |S_0| + \phi(r)\lambda \quad (4)$$

Also, by fixing the values of $S_0$ and $r$, we can optimize $\lambda$ such that

$$\lambda \leq \frac{r - |S_0|}{\phi(r)} \quad (5)$$

To obtain the greatest value of $\lambda$, one must have to maximize $\frac{r - |S_0|}{\phi(r)}$. For this purpose, after some calculations for maximization of a function, we get an equation

$$\phi(r) - (r - |S_0|)\phi'(r) = 0 \quad (6)$$

Let $R^*$ be an optimized radius of the closed ball. Clearly, it will satisfy the Eq. (6).

On the other hand, if we fix $\lambda$, then from (3) we can optimize the value of $S_0$

$$|S_0| \leq r - \phi(r)\lambda \quad (7)$$

Again, the maximum value of $r - \phi(r)\lambda$ gives the maximum $S_0$. In this case, assume that $R^{**}$ be the optimized radius which will satisfy the expression $\phi'(r) = \frac{1}{\lambda}$.

**Case 2.** Assume that $\overline{B}(r,\Theta) = \{ S \in X : \| S - S_0 \| \leq r, r > 0 \}$ be a closed ball with center at $S_0$, initial condition, and radius $r$.

Now, consider

$$\|\mathcal{H}(x,t) - S_0\| = \left\| S_0 + \int_0^t G_1(x,\omega)d\omega - S_0 \right\|,$$
where,

\[ G_1 \equiv G_1 \left( S, I_0 + \int_0^\omega \int G_2 (S, I, R, I_x, I_{xx}) (x, u) du, R_0 + \int_0^\omega \int G_3 (S, I, R, R_x, R_{xx}) (x, v) dv, S_x, S_{xx} \right) \]

\[ \Rightarrow \| H(x, t) - S_0 \| \leq \phi(r) t \leq \phi(r) \lambda. \]

For \( H \) to be a self map it should have

\[ \phi(r) \lambda \leq r. \] (8)

The fix values of \( S_0 \) and \( \lambda \) give the optimized radius, such that

\[ r \geq \phi(r) \lambda. \] (9)

Again, for the fixed values of \( S_0 \) and \( r \), we can optimize \( \lambda \) such that

\[ \lambda \leq \frac{r}{\phi(r)} \] (10)

Similarly, to obtain the large interval of continuity, one must have to maximize \( \frac{r}{\phi'(r)} \), where \( \phi(r) \) is supposed to be differentiable so, the optimal radius \( r = R^{**} \) should satisfy the equation

\[ \phi(r) = r \phi'(r) \] (11)

If, \( \phi(r) \) is twice differentiable on \( R \) then,

\[ \frac{d^2}{dr^2} \left( \frac{r}{\phi(r)} \right) = -\frac{r \phi''(r)}{(\phi(r))^2}. \]

If, \( \phi''(r) > 0 \), for all values of \( r \) then, \( \frac{r}{\phi(r)} \) will be maximum for every \( R^{**} \), which is the exact root of (11). According to the theory of calculus, between any two maxima for a function, there exists at least one point at which the function has minimum value. Therefore, with the presence of condition \( \phi''(r) > 0 \), Eq. (11) has not more than one solution. An important result can be developed in the light of above discussion. The following theorem is established in this respect.

**Theorem:** Suppose that, \( \phi(r) \) be a continuously differentiable function then, the equation \( \phi(r) = r \phi'(r) \) has a root which becomes an optimal radius. Moreover, for \( \phi(r) \in C^2 \) and \( \phi(r) > 0 \), there exists at most one optimal radius of a closed and convex subset of a Banach space \( X \) which possesses the solution of \( (1) \). Numerical analysis has a prominent role in the study of applied mathematics, specially for the analysis of epidemic models. To find the solutions of these models, we use different numerical techniques and through simulation we confirm the behavior of the solution. The numerical techniques which preserve the important properties of the model, for instance, positivity, consistency, stability, convergence and boundedness [22–24]. In the next section, a suitable nonstandard finite difference scheme is applied to obtain the numerical solution.

### 3 Numerical Analysis of Proposed Model

The finite difference methods make, the approximate solutions of the systems involving linear and nonlinear systems of partial differential equations, easy [25]. In these techniques, we convert the continuous model in to a discrete formulation choosing the finite number of function values at some finite number of points in the domain, which is easy to handle. The Taylor’s series is the bestway to obtain these approximations. Now, let \( M \) and \( N \) be any to finite positive integers and
$T$ be any positive real number. The spatial interval $[a, b]$ over the time period $[0, T]$ are discretized according to the partitions $a = x_0 < x_1 < x_2 < \ldots < x_M = b$ and $0 = t_0 < t_1 < t_2 < \ldots < t_N = T$ respectively, with the norm $h = \frac{b-a}{M}$ and $k = \frac{T}{N}$. Divide $[a, b] \times [0, T]$ having $M \times N$ grid points with space and time step sizes $h$ and $k$ respectively. The points of the partitions now become $x_i = ih$ and $t_n = nk$, where $i = 0, 1, 2, \ldots, M$ and $n = 0, 1, 2, \ldots, N$. Suppose that, $S^n_i, I^n_i$ and $R^n_i$ denote the approximations of $S(x, t), I(x, t)$ and $R(x, t)$ respectively at the grid points $(ih, nk)$. In this article, we use a non-standard finite difference implicit scheme carrying some important physical properties for the discrete model, developed in [26]. Discrete model equations form a matrix or iterative process that are used to find the best approximation of the solution to the system (1). The system (1) can be converted in to discrete form by using the following approximations.

\[
\begin{align*}
F_i &= \frac{F_{i+1}^n - F_i^n}{k} \quad F_x = \frac{F_{i+1}^n - F_{i-1}^n}{h} \quad F_{xx} = \frac{F_{i+1}^n - 2F_i^n + F_{i-1}^n}{h^2} \\
S_i^{n+1} - S_i^n + \alpha_1 \left( \frac{S_{i+1}^{n+1} - S_{i-1}^{n+1}}{h} \right) &= a - c I_i^n S_i^n (1 + \gamma I_i^n) - \mu S_i^{n+1} + \alpha R_i^n \\
&+ d_1 \left( \frac{S_{i-1}^{n+1} - 2S_i^{n+1} + S_{i+1}^{n+1}}{h^2} \right), \\
S_i^{n+1} - S_i^n + \frac{\alpha_1 k}{h} \left( S_{i+1}^{n+1} - S_{i-1}^{n+1} \right) &= ak - c k I_i^n S_i^n (1 + \gamma I_i^n) - \mu k S_i^{n+1} + \alpha k R_i^n \\
&+ \frac{d_1 k}{h^2} \left( S_{i-1}^{n+1} - 2S_i^{n+1} + S_{i+1}^{n+1} \right). \\
\end{align*}
\]  

From first Eq. of (1),
\[
S_i^{n+1} - S_i^n + g_1 \left( S_{i+1}^{n+1} - S_{i-1}^{n+1} \right) = ak - c k I_i^n S_i^n (1 + \gamma I_i^n) - \mu k S_i^{n+1} + \alpha k R_i^n \\
+ g_2 \left( S_{i-1}^{n+1} - 2S_i^{n+1} + S_{i+1}^{n+1} \right), \\
- (g_1 + g_2) S_{i-1}^{n+1} + (1 + 2g_2 + c k I_i^n (1 + \gamma I_i^n) + \mu k) S_i^{n+1} - g_2 S_{i+1}^{n+1} = ak + \alpha k R_i^n + S_i^n.
\]  

Again, second Eq. of (1) becomes as
\[
I_i^{n+1} - I_i^n + \alpha_2 \left( \frac{I_{i+1}^{n+1} - I_{i-1}^{n+1}}{h} \right) = c I_i^n S_i^n (1 + \gamma I_i^n) - (\beta + \mu + \delta - b) I_i^{n+1} + d_2 \left( \frac{I_{i-1}^{n+1} - 2I_i^{n+1} + I_{i+1}^{n+1}}{h^2} \right),
\]
Put $g_1' = \frac{a_1 k}{h}$ and $g_2' = \frac{d k}{h^2}$

$$-(g_1' + g_2') I_{i+1}^{n+1} + (1 + g_1' + k (\beta + \mu + \delta - \beta) + 2g_2') I_{i}^{n+1} - g_2' I_{i+1}^{n+1} = ck I_{i}^{n} S_{i}^{n} (1 + \gamma I_{i}^{n}) + I_{i}^{n}.$$  \hspace{1cm} (14)

Also, the third Eq. of (1) becomes as follows,

$$\frac{R_{i+1}^{n+1} - R_{i}^{n+1}}{k} + \alpha_3 \left( \frac{R_{i+1}^{n+1} - R_{i+1}^{n+1}}{h} \right) = \beta I_{i}^{n} - (\alpha + \mu) R_{i}^{n+1} + d_3 \left( \frac{R_{i-1}^{n+1} - 2R_{i}^{n+1} + R_{i+1}^{n+1}}{h^2} \right).$$

Insert $g_1'' = \frac{a_2 k}{h}$, $g_2'' = \frac{d k}{h^2}$ in above, we get

$$-(g_1'' + g_2'') R_{i-1}^{n+1} + (1 + g_1'' + k (\alpha + \mu) + 2g_2'') R_{i}^{n+1} - g_2'' R_{i+1}^{n+1} = \beta k I_{i}^{n} + R_{i}^{n}.$$  \hspace{1cm} (15)

4 Physical Properties of the Proposed Numerical Scheme

4.1 Consistency of the Discrete Model

After developing the discrete model from the continuous model, it is very important to check whether the discrete model is consistent with the given continuous model because, the solution of the discrete model approximates the exact solution associated to the continuous model. We also observe the order of consistency of our proposed scheme. After using the approximations (12) and Taylor series in all the equations of (1), we get, first equation of (1)

$$S_{i}^{n+1} + k \frac{\partial S}{\partial t} + \frac{k^2}{2!} \frac{\partial^2 S}{\partial t^2} + \frac{k^3}{3!} \frac{\partial^3 S}{\partial t^3} + \cdots - S_{i}^{n} = \frac{\alpha_1 k}{h} \left( S_{i}^{n} + k \frac{\partial S}{\partial t} + \frac{k^2}{2!} \frac{\partial^2 S}{\partial t^2} + \frac{k^3}{3!} \frac{\partial^3 S}{\partial t^3} + \cdots \right)$$

$$+ \frac{\alpha k}{h} \left( S_{i}^{n} + k \frac{\partial S}{\partial t} + \frac{k^2}{2!} \frac{\partial^2 S}{\partial t^2} + \frac{k^3}{3!} \frac{\partial^3 S}{\partial t^3} + \cdots \right)$$

$$- \frac{\alpha k}{h} \left( S_{i}^{n} + k \frac{\partial S}{\partial t} + \frac{k^2}{2!} \frac{\partial^2 S}{\partial t^2} + \frac{k^3}{3!} \frac{\partial^3 S}{\partial t^3} + \cdots \right)$$

$$\times \left( c k I_{i}^{n} (1 + \gamma I_{i}^{n}) + \mu k \right) + \alpha k R_{i}^{n} + \frac{d k}{h^2} \left( S_{i}^{n} + k \frac{\partial S}{\partial t} + \frac{k^2}{2!} \frac{\partial^2 S}{\partial t^2} + \frac{k^3}{3!} \frac{\partial^3 S}{\partial t^3} + \cdots \right)$$

$$\times \left( c k I_{i}^{n} (1 + \gamma I_{i}^{n}) + \mu k \right) + \alpha k R_{i}^{n} + \frac{d k}{h^2} \left( S_{i}^{n} + k \frac{\partial S}{\partial t} + \frac{k^2}{2!} \frac{\partial^2 S}{\partial t^2} + \frac{k^3}{3!} \frac{\partial^3 S}{\partial t^3} + \cdots \right)$$

$$- \frac{k^3}{3!} \frac{\partial^3 S}{\partial t^3} + \cdots \bigg) \bigg).$$

Taking $k$ common from both sides and apply $k \to 0$ and $h \to 0$, we get

$$\frac{\partial S}{\partial t} + \frac{\partial S}{\partial x} = a - c I S (1 + \gamma I) - \mu S + \alpha R + \frac{d_1}{h^2} S / \partial x^2.$$
4.2 Stability of the Proposed Scheme

From Eq. (13)

\[ - (g_1 + g_2) S_{n-1}^{n+1} + \left( 1 + 2g_2 + c k I_i^n \left( 1 + \gamma I_i^n \right) + \mu k \right) S_i^n + 1 - g_2 S_{i+1}^{n+1} = ak + \alpha k R_i^n + S_i^n \]

After performing the process of linearization, we get,

\[ - (g_1 + g_2) \varphi_1 (t + \Delta t) e^{i w(x - \Delta x)} + \left( 1 + 2g_2 + ck (1 + \gamma) + \mu k \right) \varphi_1 (t + \Delta t) e^{i w x} - g_2 \varphi_1 (t + \Delta t) e^{i w(x + \Delta x)} \]

\[ = \varphi_1 (t) e^{i w x} \]

\[ \frac{\varphi_1 (t + \Delta t)}{\varphi_1 (t)} = - (g_1 + g_2) e^{i w(x - \Delta x)} + \left( 1 + 2g_2 + c k (1 + \gamma) + \mu k \right) e^{i w x} - g_2 e^{i w(x + \Delta x)} \]

\[ \left| \frac{\varphi_1 (t + \Delta t)}{\varphi_1 (t)} \right| = \frac{1}{- (g_1 + g_2) e^{- \Delta w x} + \left( 1 + 2g_2 + c k (1 + \gamma) + \mu k \right) - g_2 e^{\Delta w x}} \leq 1. \]

Eq. (14) implies

\[ - (g'_1 + g'_2) I_{i-1}^{n+1} + \left( 1 + g'_1 + k (\beta + \mu + \delta - b) + 2g'_2 \right) I_i^{n+1} - g'_2 I_{i+1}^{n+1} = c k I_i^n S_i^n \left( 1 + \gamma I_i^n \right) + I_i^n \]

\[ \Rightarrow - (g'_1 + g'_2) \varphi_2 (t + \Delta t) e^{i w(x - \Delta x)} + \left( 1 + g'_1 + k (\beta + \mu + \delta - b) + 2g'_2 \right) \varphi_2 (t + \Delta t) \]

\[ \times e^{i w x} - g'_2 \varphi_2 (t + \Delta t) e^{i w(x + \Delta x)} = \varphi_2 (t) e^{i w x} \]

This implies that

\[ \left| \frac{\varphi_2 (t + \Delta t)}{\varphi_2 (t)} \right| = \frac{1}{- (g_1 + g_2) e^{- \Delta w x} + \left( 1 + g_1 + k (\beta + \mu + \delta - b) + 2g_2 \right) - g_2 e^{\Delta w x}} \leq 1. \]
Similarly, from Eq. (15), we also have
\[
\left| \frac{\varphi_3(t + \Delta t)}{\varphi_3(t)} \right| \leq 1.
\]
which shows that the proposed scheme is Von-Neumann stable.

4.3 Positivity

According to [27], M-matrix theory is a reliable tool for the verification of the positivity for most of the numerical models concerning engineering, economics, autocatalytic chemical reactions etc. A square matrix over a real field \( R \) is said to be an \( M \)-matrix if all entries in the off-diagonal are non-positive.

4.4 Z-Matrix

If all the off-diagonal entries of a real matrix \( A \) are non-positive then \( A \) is called a \( Z \)-matrix.

4.5 M-Matrix

A square matrix \( A \) over \( R \) is an \( M \)-matrix if:

(i) \( A \) is a \( Z \)-matrix.
(ii) all the diagonal components of \( A \) are positive,
(iii) \( A \) is strictly diagonally dominant.

It is important to note that \( M \)-matrices are non-singular so are invertible and their inverses are positive matrices [28]. This important property will be used in the next result.

Remark: It is clear that \( M \)-matrix is a diagonally dominant and its inverse always consists of positive entries.

Theorem: For any \( h \geq 0 \) and \( k \geq 0 \) the discretized system (13)–(15) has positive solutions. That is, \( S^n \geq 0 \), \( I^n \geq 0 \) and \( R \geq 0 \) for all \( n = 0, 1, 2, 3, \ldots, N \).

Proof: It is important to note that the discrete system (13)–(15) generated from finite difference method can be converted in the vector form, such as

\[ DS^{n+1} = ak + akR^n_i + S^n_i. \]  
(16)

\[ EI^{n+1} = ckl^n_iS^n_i(1 + \gamma I^n_i) + I^n_i. \]  
(17)

\[ FR^{n+1} = \beta kI^n_i + R^n_i. \]  
(18)

In the above setting, \( D, E, F \) are square matrices of order \( (M + 1) \).

Then
\[
D = \begin{pmatrix}
(\beta_1)^n_0 & \beta_2^n & 0 & 0 & 0 & 0 \\
\beta_3^n & (\beta_1)^n_1 & \beta_4^n & \cdots & 0 & 0 \\
0 & \beta_3^n & (\beta_1)^n_2 & \cdots & 0 & 0 \\
\vdots & \vdots & \ddots & \ddots & \vdots \\
0 & 0 & 0 & (\beta_1)^n_{M-2} & \beta_4^n & 0 \\
0 & 0 & 0 & \cdots & \beta_3^n & (\beta_1)^n_{M-1} \\
0 & 0 & 0 & \cdots & \beta_3^n & (\beta_1)^n_M
\end{pmatrix}
\]
That is, the proposed numerical scheme preserves the positivity of the solution.

\( E = \begin{pmatrix} (\gamma_1^n)^n_0 & \gamma_2 & 0 & 0 & 0 & 0 \\ \gamma_3 & (\gamma_1^n)_1 & \gamma_4 & \cdots & 0 & 0 \\ 0 & \gamma_3 & (\gamma_1^n)_2 & 0 & 0 & 0 \\ \vdots & \vdots & \vdots & \ddots & \vdots & \vdots \\ 0 & 0 & 0 & \cdots & \gamma_3 & (\gamma_1^n)_{M-2} \\ 0 & 0 & 0 & \cdots & \gamma_3 & (\gamma_1^n)_{M-1} \\ 0 & 0 & 0 & 0 & \gamma_3 & (\gamma_1^n)_M \end{pmatrix} \)

\( F = \begin{pmatrix} (\delta_1^n)_0 & \delta_2 & 0 & 0 & 0 & 0 \\ \delta_3 & (\delta_1^n)_1 & \delta_4 & \cdots & 0 & 0 \\ 0 & \delta_3 & (\delta_1^n)_2 & 0 & 0 & 0 \\ \vdots & \vdots & \vdots & \ddots & \vdots & \vdots \\ 0 & 0 & 0 & \cdots & \delta_3 & (\delta_1^n)_{M-2} \\ 0 & 0 & 0 & \cdots & \delta_3 & (\delta_1^n)_{M-1} \\ 0 & 0 & 0 & 0 & \delta_3 & (\delta_1^n)_M \end{pmatrix} \)

The diagonal elements of \( \mathcal{A}^* \) are \((\beta_1^n)^n_0 = 1 + \mu k + 2g_2 + c k I^n_0 I^n (1 + \gamma I^n_0), (\beta_1^n)_r = 1 + \mu k + c k I^n_0 (1 + \gamma I^n_0), r = 0, 1, 2, \ldots, M \) and for all \( n \in \{0, 1, 2, \ldots, N\} \) and the values of off-diagonal elements are \( \beta_2 = -(g_1 + 2g_2), \beta_3 = -(g_1 + g_2), \beta_4 = -g_2. \)

The diagonal elements of \( \mathcal{Y} \) are \((\gamma_1^n)^n_0 = 1 + g_1' + k(\beta + \mu + \delta - b) + 2g_2', (\gamma_1^n)_r = 1 + g_1' + k(\beta + \mu + \delta - b) + g_2', r = 0, 1, 2, \ldots, M \) and for all \( n \in \{0, 1, 2, \ldots, N\} \) and the values of off-diagonal elements are \( \gamma_2 = -(g_1' + 2g_2'), \gamma_3 = -(g_1' + g_2'), \gamma_4 = -g_2'. \)

Similarly, the diagonal and off-diagonal entries of \( \mathcal{Z} \) are, \((\delta_1^n)_0 = 1 + k(\alpha + \mu) + 2g_2'', (\delta_1^n)_r = 1 + k(\alpha + \mu) + g_2'', r = 0, 1, 2, \ldots, M; \) for all \( n \in \{0, 1, 2, \ldots, N\} \) and \( \delta_2 = -(g_1'' + 2g_2''), \delta_3 = -(g_1'' + g_2''), \delta_4 = -g_2'' \) respectively.

The entries in the column matrices of the system (16)–(18) are \( ak + ak R^n_0 + S^n_1, c k I^n_0 S^n_1 + I^n_0, b k I^n_0 R^n_0 \) respectively. Also, since \( S^n = (S^n_0, S^n_1, \ldots, S^n_M)^T \) and \( I^n = (I^n_0, I^n_1, \ldots, I^n_M)^T, R^n = (R^n_0, R^n_1, \ldots, R^n_M)^T \) and \( S^n_0, I^n_0 \) and \( R^n_0 \) are non-negative so are the diagonal entries of \( D, E \) and \( F. \) Moreover, all the off-diagonal entries of the matrices \( D, E, F \) are negative and the matrices \( D, E, F \) are strictly diagonally dominant. This implies that the matrices \( D, E \) and \( F \) are M-matrices which concludes that \( D, E \) and \( F \) are non-singular and hence invertible. So (16)–(18) can be written as

\[ S^{n+1} = D^{-1}(ak + ak R^n_0 + S^n_1). \]

\[ I^{n+1} = E^{-1}(ck I^n_0 S^n_1 (1 + \gamma I^n_0) + I^n_0) \]

\[ R^{n+1} = F^{-1}(b k I^n_0 R^n_0) \]

Now, suppose that \( S^n > 0, I^n > 0 \) and \( R^n > 0 \) and \( \mathcal{X}, \mathcal{Y}, \mathcal{Z} \) satisfy all the conditions on M-matrix which concludes that all the entries of the matrices \( D^{-1}, E^{-1}, F^{-1} \) are positive. So, \( S^{n+1} > 0, I^{n+1} > 0 \) and \( R^{n+1} > 0. \) Hence, by induction, the system (16)–(18) has a positive solution. That is, the proposed numerical scheme preserves the positivity of the solution.
5 Numerical Example and Simulations

In this experiment, the following initial conditions are supposed

\[ S(x,0) = \begin{cases} 
0.5x, & 0 \leq x < 0.5 \\
0.5(1-x), & 0.5 \leq x \leq 1
\end{cases} \]

\[ I(x,0) = \begin{cases} 
0.4x, & 0 \leq x < 0.5 \\
0.4(1-x), & 0.5 \leq x \leq 1
\end{cases} \]

Figure 1: (a) The spatio-temporal simulations results for susceptible population, (b) the spatio-temporal simulations results for infected population, (c) the spatio-temporal simulations results for recovered population, (d) the graphical simulations in 2-D plot for all sub-populations by considering \( x = 1 \)
\[ R(x,0) = \begin{cases} 
0.1x, & 0 \leq x < 0.5 \\
0.1(1-x), & 0.5 \leq x \leq 1 
\end{cases} \]

### 5.1 Disease Free State

For the disease free state we take the following values:

\[
a = 0.5, \quad \mu = 0.5, \quad \delta = 0.05, \quad b = 0.205, \quad \beta = 0.09871, \quad c = 0.380, \quad \gamma = 0.0003, \quad \alpha = 0.854302.
\]

**Figure 2:** (a) The spatio-temporal simulations results for susceptible population, (b) the spatio-temporal simulations results for infected population, (c) the spatio-temporal simulations results for recovered population, (d) the graphical simulations in 2-D plot for all sub-populations by considering \( x = 1 \)
In the Fig. 1, we take the values of parameters in such a way that the value of reproductive number is less than 1. The mesh graphs and combined 2-D plot of all sub-population model are provided in Fig. 1. The SIR corona virus advection diffusion model is manifested two steady states, virus free state and virus existence state. Since in Fig. 1, the value of reproductive number is less than unity therefore model (1) demonstrates the state when virus in not present in the population. Also we see that the state variables present in the system (1) are the population densities. The graphical solutions depicted in the Fig. 1 indicate that the designed technique sustains the positive solution of model (1) and validates the theorem of positivity. Also the stability of virus free steady state of continuous model (1) is retained by designed technique which is again demonstrated in Fig. 1.

5.2 Endemic State

For the endemic state we take the following values:

\[ a = 0.5, \mu = 0.5, \delta = 0.05, b = 0.205, \beta = 0.09871, c = 0.580, \gamma = 0.0003, \alpha = 0.854302. \]

Fig. 2 indicates the solution mesh graphs and combined two dimensional plot for the steady state at which the virus exists in the population by using proposed technique. Again the positive solution of system (1) is sustained by the upwind NSFD technique. As the values of parameters describe that the value of reproductive quantity is greater than 1, so the corona virus is present in the population and the advection diffusion system (1) possesses the virus persistent in the population. The graphical behavior of all state variables by using proposed technique in the Fig. 2 depicts that the virus is persisted in the population. This shows that the underlying technique is consistent with the continuous model (1).

6 Conclusion

In the current article, we have analyzed a mathematical model concerned with the novel coronavirus disease and the existence and uniqueness of its solution in a closed and convex subset of a banach space is also discussed. The prominent feature of our proposed model is the combination of advection and diffusion in the model which makes the model more realistic and comprehensive. Numerical solution of the proposed model is also computed with the help of nonstandard finite difference scheme which preserves the structural properties of the proposed continuous model. To support the validity of our numerical scheme, we have checked some of its physical properties such as consistency, stability and positivity which is the most important property in any population dynamical model. For positivity, we have used the M-matrix theory. Numerical simulations play a vital role in the study of numerical analysis so, to check the graphical behavior of solution, numerical simulation is also performed that verifies our results. We can extend our work to two and three dimensions.

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