Research Article

Numerical Solution of Fractional-Order HIV Model Using Homotopy Method

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In this study, we construct a convergent algorithm for generating an approximate analytic solution for the fractional HIV infection of CD4+ T cells with Atangana–Baleanu fractional derivatives in the Caputo sense. We compute the solution by utilizing the fractional homotopy analysis transform method (FHATM) and achieved a convergence region of the solution by employing an auxiliary parameter. Moreover, we apply a numerical scheme proposed by Toufik and Atangana for solving this kind of problem and compared with our results. A good agreement between the new algorithm and the numerical scheme is remarkable. The solution via the present algorithm can be obtained without any linearization or discretization which makes it reliable and easy to apply.

1. Introduction

Fractional calculus has played a significant role within the field of science and engineering, and many mathematicians and scientists have been working in this field lately. In recent decades, fractional calculus has been used in several areas of physics, biology, engineering, and others. Further details about fractional calculus and its applications can be found in the literature [1–9].

Because most nonlinear fractional differential equations cannot be solved exactly, it is necessary to use approximate and numerical methods. Various powerful mathematical techniques such as the Adomian decomposition method (ADM) [10, 11], homotopy analysis method (HAM) [12–15], optimal homotopy asymptotic method (OHAM) [16], homotopy perturbation method (HPM) [17], and variational iterative method (VIM) [18, 19] have been used to obtain an exact and approximate analytical solution.

The HAM was first introduced and employed in 1992 by Liao [20], after which many researchers successfully applied this method to solve linear and nonlinear differential equations. In recent years, many researchers have devoted their attention to obtaining a solution of linear and nonlinear differential equations using a variety of methods based on Laplace transform such as the Laplace decomposition method (LDM) [21] and the homotopy perturbation transform method (HPTM) [22]. Khan et al. [23] and Kumar et al. [24–26] coupled the HAM with Laplace transform to solve a nonlinear differential equation and Volterra integral equation. The homotopy analysis transform method (HATM) is a combination of HAM with Laplace transformation. The main advantage of this method is its ability to combine two powerful methods to obtain a rapid convergent series for fractional differential equations. This method provides us with a convenient way to control the convergence of the series solution.

Recently, Toufik and Atangana [27] developed a numerical scheme to solve a nonlinear fractional differential equation considering the Atangana–Baleanu fractional derivative. This method is a combination of the fundamental fractional calculus theorem with two-step Lagrange polynomial which is successfully used to solve many real-world problems [28–30].

Since the early 1980s, researchers have made an enormous effort to mathematically model the human immunodeficiency virus (HIV), the virus responsible for causing acquired immune deficiency syndrome (AIDS). In 1989,
Perelson [31] considered the interaction of uninfected (T) and infected CD4+ (I) T cells, and free virus molecules (F) in his model, following which Perelson et al. [32] extended the original model [31]. Culshaw and Ruan [33] reduced the model discussed in [32] as follows:

\[
\frac{dT}{dt} = p - \mu_T T + kT \left(1 - \frac{T + I}{T_{\text{max}}} \right) - k_1 FT, \quad \text{(1)}
\]

\[
\frac{dI}{dt} = k_1 FT - \mu_I I, \quad \text{(2)}
\]

\[
\frac{dF}{dt} = M\mu_0 I - k_1 FT - \mu_F F, \quad \text{(3)}
\]

where \(T(t)\), \(I(t)\), and \(F(t)\) represent the concentration of healthy CD4+ T cells at time \(t\), infected CD4+ T cells, and the free HIV virus at time \(t\), respectively. Table 1 summarizes the meanings of functions and parameters. Equation (1) describes the rate of change in the uninfected population of CD4+ T cells. The first term is the constant rate at which the body produces CD4+ T cells from precursors in the bone marrow. Because the virus can infect both thymocytes and T cells, as for all cells in the body, these cells have a finite lifetime; thus, the second term describes the decreasing source. The third term describes the logistic growth of the healthy CD4+ T cells, and the proliferation of infected CD4+ T cells is neglected. The last term models the rate at which the free virus infects a CD4+ T cell. Once a T cell has been infected, it becomes an infected cell; therefore, \(k_1\) FT is subtracted from equations (1) and (3) and added to equation (2). Hence, \(F\) and \(T\) decrease concurrently.

Equation (2) describes the rate of change in the infected population of actively infected T cells. The first term represents the rate of infection of CD4+ T cells by the virus. The second term represents the rate of disappearance of infected cells.

The three terms in equation (3) refer to the rate of production and destruction of the free infection virus. An actively infected CD4+ T cell produces \(M\) virus particles; thus, the rate at which the virus is produced is set equal to \(M\) times the lytic death rate for the infected cell. A free virus is lost as a result of binding to an uninfected CD4+ T cell at \(k_1\) FT. The third term accounts for the loss of viral infectivity, viral death, and/or clearance from the body.

In this study, our approach to solving the fractional HIV model is to determine the order in which the fractional derivative changes by extending the classical HIV model (1)–(3) to the following set of fractional ordinary differential equations of the order \(\alpha\), \(\beta\), and \(\gamma\):

\[
\begin{align*}
\ABC{\alpha}{D}_t^\alpha T(t) &= p - \mu_T T + kT \left(1 - \frac{T + I}{T_{\text{max}}} \right) - k_1 FT, \quad 0 < \alpha < 1, \\
\ABC{\beta}{D}_t^\beta F(t) &= k_1 FT - \mu_I I, \quad 0 < \beta < 1,
\end{align*}
\]

where \(\ABC{\alpha}{D}_t^\alpha f(t)\), \(\ABC{\beta}{D}_t^\beta f(t)\), and \(\ABC{\gamma}{D}_t^\gamma f(t)\) are the Atangana–Baleanu fractional derivative in the Caputo sense (ABC). To the best of our knowledge, this is the first work that solves fractional HIV infection of the CD4+ T cells model in ABC sense analytically and numerically. To indicate the strength of our proposed method, we compare our findings with the algorithm of Toufik and Atangana [27].

**2. Preliminaries and Notations**

The Atangana–Baleanu fractional derivative in the Caputo sense (ABC) is defined as [6, 34]

\[
\ABC{\alpha}{D}_t^\alpha f(t) = \frac{M(\alpha)}{1 - \alpha} \int_0^t \frac{d}{ds} f(s) \mathcal{E}_\alpha \left( -\frac{\alpha}{1 - \alpha} (t - s) \right) ds, \quad n - 1 < \alpha \leq n,
\]

where \(\alpha \in \mathbb{R}, M(\alpha) > 0\) is a normalization function satisfying

\[
M(\alpha) = (1 - \alpha) + \frac{\alpha}{\Gamma(\alpha)},
\]

with \(M(0) = M(1) = 1\), \(\mathcal{E}_\alpha(.)\) denotes the Mittag-Leffler function, defined by

Table 1: List of parameters and functions.

| Parameters and functions | Description | Values |
|--------------------------|-------------|--------|
| \(T(t)\)                | Concentration of uninfected CD4+ T cells | \(T(0) = 1000\) |
| \(I(t)\)                | Concentration of infected CD4+ T cells | \(I(0) = 0\) |
| \(F(t)\)                | Concentration of HIV RNA | \(F(0) = 0.001\) |
| \(\mu_T\)               | Natural death rate of CD4+ T cells (concentration) | 0.02 |
| \(\mu_I\)               | Blanket death rate of infected CD4+ T cells | 0.26 |
| \(\mu_F\)               | Lytic death rate of infected cells | 0.24 |
| \(k_1\)                 | Rate at which CD4+ T cells become infected with the virus | 2.4 \times 10^{-5} |
| \(k_1\)                 | Rate at which infected cells become active | 2 \times 10^{-5} |
| \(k\)                   | Growth rate of concentration of CD4+ T cells | 0.03 |
| \(M\)                   | Number of virion produced by infected CD4+ T cells | 500 |
| \(T_{\text{max}}\)      | Maximal concentration of CD4+ T cells | 1500 |
| \(p\)                   | Source term for uninfected CD4+ T cells | 10 |

\[
F(t) = M\mu_0 I - k_1 FT - \mu_F F, \quad 0 < \gamma < 1,
\]
and \( \Gamma (z) = \int_0^\infty t^{z-1} e^{-t} \, dt \), \( \Re (z) \).

The fractional integral for the ABC, which is newly defined with a nonlocal kernel and does not have singularities at \( t = s \), is defined as follows [6]:

\[
\mathcal{L}_a^{ABC} I_t^a f(t) = \frac{1 - \alpha}{M(\alpha)} f(t) + \frac{\alpha}{M(\alpha) \Gamma (\alpha)} \int_a^t f(s)(t-s)^{\alpha-1} \, ds.
\]

Here, when \( \alpha \) equals zero, the initial function is recovered, and when \( \alpha \) equals unity, the classical ordinary integral is obtained.

The Laplace transform of the fractional definitions with ABC is given as follows [6]:

\[
\mathcal{L}_a^{ABC} I_t^a f(t) \bigg| (s) = \left( \frac{M(\alpha) s^\alpha \mathcal{L} f(t)(s) - s^{\alpha-1} f(\alpha)}{1 - \alpha} \right) \frac{\alpha}{(\alpha/1-\alpha)}.
\]

3. Homotopy and Laplace Transform for FHATM

Applying the Laplace transform to equations (4)–(6) and using the formula for the Laplace transform of the ABC and then simplifying these equations, we find that

\[
\mathcal{L} \left[ T(t; q) \right] = \frac{T(0)}{s} + \frac{p}{s} + \frac{s^{\alpha}(1-\alpha) + \alpha}{s^\alpha M(\alpha)} - \frac{\alpha}{s^{\alpha} M(\alpha)} \mathcal{L}
\]

\[
\left\{ \left( (k - \mu_2)T(t; q) - \frac{k}{T_{\max}} (T(t))^2 - \frac{k}{T_{\max}} T \right) \right\},
\]

\[
\mathcal{L} \left[ I(t; q) \right] = \frac{I(0)}{s} + \frac{\beta}{s^\beta M(\beta)} \mathcal{L} \left[ k_1^\prime F(t)(T(t) - \mu_1 I(t)) \right],
\]

\[
\mathcal{L} \left[ F(t; q) \right] = \frac{F(0)}{s} + \frac{s^{\gamma}(1-\gamma) + \gamma}{s^\gamma M(\gamma)} \mathcal{L} \left[ k_2^\prime \mu_2 \phi_1(t; q) - \mu_2 \phi_2(t; q) \right]
\]

Next, defining the nonlinear operators as

\[
N_T[\phi_1(t; q), \phi_2(t; q), \phi_3(t; q)] = \mathcal{L}[\phi_1(t; q)] - \frac{T(0)}{s} \frac{p}{s} \frac{s^\alpha(1-\alpha) - \alpha}{s^\alpha M(\alpha)} - \frac{s^\alpha(1-\alpha) + \alpha}{s^\alpha M(\alpha)} \mathcal{L}
\]

\[
\mathcal{L} \left[ k_1^\prime [\phi_2(t; q) - \phi_3(t; q)] \right],
\]

\[
N_I[\phi_1(t; q), \phi_2(t; q), \phi_3(t; q)] = \mathcal{L}[\phi_2(t; q)] - \frac{I(0)}{s} - \frac{s^{\beta}(1-\beta) + \beta}{s^\beta M(\beta)} \mathcal{L} \left[ k_1^\prime \phi_3(t; q) \phi_1(t; q) - \mu_2 \phi_2(t; q) \right],
\]

\[
N_F[\phi_1(t; q), \phi_2(t; q), \phi_3(t; q)] = \mathcal{L}[\phi_3(t; q)] - \frac{F(0)}{s} - \frac{s^{\gamma}(1-\gamma) + \gamma}{s^\gamma M(\gamma)} \mathcal{L} \left[ k_2^\prime \mu_2 \phi_1(t; q) - k_3 \phi_3(t; q) \phi_1(t; q) - \mu_2 \phi_3(t; q) \right],
\]

where \( N_T, N_I, \) and \( N_F \) are the nonlinear operators. Let \( h \) be a nonzero auxiliary parameter. Using the embedding parameter \( q \in [0, 1] \), we construct the so-called zeroth-order deformation equation:

\[
(1-q) \mathcal{L} \left[ \phi_1(t; q) - T_0(t) \right] = q h N_T[\phi_1(t; q), \phi_2(t; q), \phi_3(t; q)],
\]

\[
(1-q) \mathcal{L} \left[ \phi_2(t; q) - I_0(t) \right] = q h N_I[\phi_1(t; q), \phi_2(t; q), \phi_3(t; q)],
\]

\[
(1-q) \mathcal{L} \left[ \phi_3(t; q) - F_0(t) \right] = q h N_F[\phi_1(t; q), \phi_2(t; q), \phi_3(t; q)],
\]

where \( \mathcal{L} \) is the Laplace operator, subject to the initial conditions

\[
\phi_1(0; q) = T_0(0), \quad \phi_2(0; q) = I_0(0), \quad \phi_3(0; q) = F_0(0).
\]

Clearly if \( q = 0 \) and \( q = 1 \) we obtain

\[
\phi_1(t; 0) = T_0(t), \quad \phi_1(t; 1) = T(t), \quad \phi_2(t; 0) = I_0(t), \quad \phi_2(t; 1) = I(t), \quad \phi_3(t; 0) = F_0(t), \quad \phi_3(t; 1) = F(t),
\]

when \( q \) varies from zero to unity, the solution of the model (4)–(6) will vary from the initial guesses \( T_0(t), I_0(t), \) and
with respect to the embedding parameter \( q \). Expanding \( \varphi_i(t; q), i = 1, 2, 3 \) by the Taylor series with respect to the embedding parameter \( q \) yields

\[
\varphi_1(t; q) = T_0(t) + \sum_{m=1}^{\infty} T_m(t)q^m,
\]

(21)

\[
\varphi_2(t; q) = I_0(t) + \sum_{m=1}^{\infty} I_m(t)q^m,
\]

(22)

\[
\varphi_3(t; q) = F_0(t) + \sum_{m=1}^{\infty} F_m(t)q^m,
\]

(23)

where

\[
T_m(t) = \frac{1}{m!} \frac{\partial^m \varphi_1(t; q)}{\partial q^m} \bigg|_{q=0},
\]

(24)

\[
I_m(t) = \frac{1}{m!} \frac{\partial^m \varphi_2(t; q)}{\partial q^m} \bigg|_{q=0},
\]

(25)

\[
F_m(t) = \frac{1}{m!} \frac{\partial^m \varphi_3(t; q)}{\partial q^m} \bigg|_{q=0}.
\]

The convergence of equations (21)–(23) depends on the nonzero auxiliary parameters \( h \) [20]. Moreover, if the initial values guessed for \( T_0(t), I_0(t), \) and \( F_0(t) \) and the auxiliary parameter \( h \) are appropriately selected, then at \( q = 1 \), series (21)–(23) converges

\[
\varphi_1(t; 1) = T_0(t) + \sum_{m=1}^{\infty} T_m(t) \text{ i.e. } T(t) = T_0(t) + \sum_{m=1}^{\infty} T_m(t),
\]

(26)

\[
\varphi_2(t; 1) = I_0(t) + \sum_{m=1}^{\infty} I_m(t) \text{ i.e. } I(t) = I_0(t) + \sum_{m=1}^{\infty} I_m(t),
\]

(27)

\[
\varphi_3(t; 1) = F_0(t) + \sum_{m=1}^{\infty} F_m(t) \text{ i.e. } F(t) = F_0(t) + \sum_{m=1}^{\infty} F_m(t),
\]

which must be one of the solution of model (4)–(6), as proved by [20]. The equations governing the unknown functions can be deduced from the zeroth-deformation equations (16)–(18). Define the vectors

\[
\overrightarrow{T}_m(t) = [T_0(t), T_1(t), \ldots, T_m(t)], \quad m = 1, 2, \ldots, n,
\]

(28)

\[
\overrightarrow{I}_m(t) = [I_0(t), I_1(t), \ldots, I_m(t)], \quad m = 1, 2, \ldots, n,
\]

\[
\overrightarrow{F}_m(t) = [F_0(t), F_1(t), \ldots, F_m(t)], \quad m = 1, 2, \ldots, n.
\]

Differentiating the zeroth-deformation equations (16)–(18) \( m \)-times with respect to the embedding parameter \( q \), then setting \( q = 0 \), and finally dividing them by \( m! \), enables the \( m \)-th-order deformation equations to be obtained:

\[
\mathcal{L}[T_m(t) - \chi_m T_{m-1}(t)] = hR_{m,T}(\overrightarrow{T}_{m-1}, \overrightarrow{T}_{m-1}, \overrightarrow{T}_{m-1}),
\]

\[
m = 1, 2, \ldots, n,
\]

(29)

\[
\mathcal{L}[I_m(t) - \chi_m I_{m-1}(t)] = hR_{m,I}(\overrightarrow{T}_{m-1}, \overrightarrow{T}_{m-1}, \overrightarrow{T}_{m-1}),
\]

\[
m = 1, 2, \ldots, n,
\]

(30)

\[
\mathcal{L}[F_m(t) - \chi_m F_{m-1}(t)] = hR_{m,F}(\overrightarrow{T}_{m-1}, \overrightarrow{T}_{m-1}, \overrightarrow{T}_{m-1}),
\]

\[
m = 1, 2, \ldots, n,
\]

(31)

with initial conditions

\[
T_m(0) = 0,
\]

(32)

\[
I_m(0) = 0,
\]

(33)

\[
F_m(0) = 0.
\]

It should be emphasized that the \( m \)-th-order deformation equations (29)–(31) are linear; hence, they can be solved by Mathematica or MATLAB. For simplicity, we can specify the auxiliary functions to be equal to unity. Applying the inverse Laplace transform to equations (29)–(31), we obtain

\[
T_m(t) = \chi_m T_{m-1}(t) + h \mathcal{L}^{-1}(R_{m,T}(\overrightarrow{T}_{m-1}, \overrightarrow{T}_{m-1}, \overrightarrow{T}_{m-1})), \quad m = 1, 2, \ldots, n,
\]

(34)

\[
I_m(t) = \chi_m I_{m-1}(t) + h \mathcal{L}^{-1}(R_{m,I}(\overrightarrow{T}_{m-1}, \overrightarrow{T}_{m-1}, \overrightarrow{T}_{m-1})), \quad m = 1, 2, \ldots, n,
\]

(35)

\[
F_m(t) = \chi_m F_{m-1}(t) + h \mathcal{L}^{-1}(R_{m,F}(\overrightarrow{T}_{m-1}, \overrightarrow{T}_{m-1}, \overrightarrow{T}_{m-1})), \quad m = 1, 2, \ldots, n,
\]

(36)
Taking initial conditions and equations (35)–(37), we obtain

\[
T_m(t) = (\chi_m + h)T_{m-1}(t) - h(1 - \chi_m)\left(T_0 + \frac{sat^a}{M(\alpha)} + \frac{s(1 - \alpha)}{M(\alpha)}\right) - h\mathcal{D}^{-1}s^a(1 - \alpha) + \alpha \mathcal{D}
\]
\[
\cdot \left\{ \left(\mu_T - k\right)T_{m-1}(t) + \frac{k}{T_{\text{max}}} \sum_{i=0}^{m-1} T_i(t)T_{m-1-i}(t) + \frac{k}{T_{\text{max}}} \sum_{i=0}^{m-1} T_i(t)I_{m-1-i}(t) + k_1 \sum_{i=0}^{m-1} T_i(t)F_{m-1-i}(t) \right\},
\]
\[
I_m(t) = (\chi_m + h)I_{m-1}(t) - h(1 - \chi_m)I_0 - h\mathcal{D}^{-1}\left\{ \frac{s^\beta(1 - \beta) + \beta}{s^\beta M(\beta)} \mathcal{D} \left[ k_1 \sum_{i=0}^{m-1} F_i(t)T_{m-1-i}(t) - \mu_I I_{m-1}(t) \right] \right\},
\]
\[
F_m(t) = (\chi_m + h)F_{m-1}(t) - h(1 - \chi_m)F_0 - h\mathcal{D}^{-1}\left\{ \frac{s^\gamma(1 - \gamma) + \gamma}{s^\gamma M(\gamma)} \mathcal{D} \left[ k_1 \sum_{i=0}^{m-1} F_i(t)T_{m-1-i}(t) + \mu_F F_{m-1}(t) - M\mu_I I_{m-1}(t) \right] \right\}.
\]

Taking initial conditions and equations (35)–(37), we obtain

\[
T_1 = -\frac{h}{M(\alpha)} \left( p + (k - \mu_T)T_0 + kT_0 \left( 1 - \frac{T_0 + I_0}{T_{\text{max}}} \right) - k_1 T_0 F_0 \right) \left( 1 - \alpha + \frac{at^a}{\Gamma(1 + \alpha)} \right),
\]
\[
I_1 = -\frac{h(kI_0T_0 - \mu_I I_0)}{M(\beta)} \left( 1 - \beta + \frac{\beta t^\beta}{\Gamma(1 + \beta)} \right),
\]
\[
F_1 = -\frac{h(M\mu_I I_0 - k_1 T_0 F_0 - \mu_F F_0)}{M(\gamma)} \left( 1 - \gamma + \frac{\gamma t^\gamma}{\Gamma(1 + \gamma)} \right),
\]
\[ T_2 = (1 + h)T_1 \]
\[ + \frac{h^2 m_i}{M(\alpha)^2} \left( k - \mu_T - \frac{k}{T_{max}} (2T_0 + I_0) - k_1 F_0 \right) \left( p + (k - \mu_T) T_0 + kT_0 \left( 1 - \frac{T_0 + I_0}{T_{max}} \right) - k_1 T_0 F_0 \right) \]
\[ - \frac{h^2 kT_0 m_2}{M(\alpha)M(\beta)} \left( k'T_0 F_0 - \mu_T I_0 \right) \]
\[ - \frac{h^2 k_1 T_0 m_3}{M(\alpha)M(\gamma)} \left( M\mu_0 I_0 - k_1 T_0 F_0 - \mu_T F_0 \right), \]

\[ I_2 = (1 + h)I_1 \]
\[ - \frac{h^2 \mu_4 m_4}{(M(\beta))^2} \left( k'I_0 T_0 - \mu_T I_0 \right) \]
\[ + \frac{h^2 k_1 F_0 m_2}{M(\alpha)M(\beta)} \left( p + (k - \mu_T) T_0 + kT_0 \left( 1 - \frac{T_0 + I_0}{T_{max}} \right) - k_1 T_0 F_0 \right) \]
\[ + \frac{h^2 k_1 T_0 m_3}{M(\gamma)(\beta)} \left( M\mu_0 I_0 - k_1 T_0 F_0 - \mu_T F_0 \right), \]

\[ F_2 = (1 + h)F_1 \]
\[ - \frac{h^2 (k_1 T_0 + \mu_T) m_0}{(M(\gamma))^2} \left( M\mu_0 I_0 - k_1 T_0 F_0 - \mu_T F_0 \right) \]
\[ - \frac{h^2 k_1 F_0 m_3}{M(\alpha)M(\gamma)} \left( p + (k - \mu_T) T_0 + kT_0 \left( 1 - \frac{T_0 + I_0}{T_{max}} \right) - k_1 T_0 F_0 \right) \]
\[ + \frac{h^2 M\mu_5 m_5}{M(\gamma)(\beta)} \left( k'I_0 T_0 - \mu_T I_0 \right), \]

where \( m_i, i = 1, 2, \ldots, 6, \) are given by

\[
\begin{align*}
m_1 &= (1 - \alpha)^2 + \frac{2(1 - \alpha)at^a}{\Gamma(1 + a)} + \frac{(at^a)^2}{\Gamma(2a + 1)}, \\
m_4 &= (1 - \beta)^2 + \frac{2(1 - \beta)\beta t^\beta}{\Gamma(1 + \beta)} + \frac{(\beta t^\beta)^2}{\Gamma(2\beta + 1)}, \\
m_6 &= (1 - \gamma)^2 + \frac{2(1 - \gamma)\gamma t^\gamma}{\Gamma(1 + \gamma)} + \frac{(\gamma t^\gamma)^2}{\Gamma(2\gamma + 1)}, \\
m_2 &= (1 - \alpha)(1 - \beta) + \frac{(1 - \alpha)\beta t^\beta}{\Gamma(1 + \beta)} + \frac{(1 - \beta)at^a}{\Gamma(1 + a)} + \frac{a\beta t^{a+\beta}}{\Gamma(a + \beta + 1)}, \\
m_5 &= (1 - \alpha)(1 - \gamma) + \frac{(1 - \alpha)\gamma t^\gamma}{\Gamma(1 + \gamma)} + \frac{(1 - \gamma)at^a}{\Gamma(1 + a)} + \frac{a\gamma t^{a+\gamma}}{\Gamma(a + \gamma + 1)}, \\
m_3 &= (1 - \beta)(1 - \gamma) + \frac{(1 - \beta)\gamma t^\gamma}{\Gamma(1 + \gamma)} + \frac{(1 - \gamma)\beta t^\beta}{\Gamma(1 + \beta)} + \frac{\beta \gamma t^{\beta+\gamma}}{\Gamma(\beta + \gamma + 1)}.
\end{align*}
\]
In a similar way, \( T_m, I_m, \) and \( F_m, \) for \( m \geq 3 \) can be obtained. Finally, the solution of model (2) is given by

\[
T(t) = \sum_{m=0}^{n-1} T_m(t), \quad (40)
\]

\[
I(t) = \sum_{m=0}^{n-1} I_m(t), \quad (41)
\]

\[
F(t) = \sum_{m=0}^{n-1} F_m(t), \quad (42)
\]

and by choosing a suitable value for \( h \) for the convergence of the series according to [20]. The analysis of the convergence of the HATM can found in the literature [35].

4. Numerical Illustration

According to [36], it should be noted that the solution of the series contain the auxiliary parameter \( h, \) which offers an easy way to control the convergence of the solution of the series. Because it is essential to assure that the series equations (25)–(27) is convergent, we plotted the \( h \) curve of 6 terms of the FHATM solution for the fractional-time ABC equations in Figures 1–3. Using these \( h \) curves, we note that the straight line that parallels the \( h \) axis provides the region of convergence. These valid regions are listed in Table 2.

Furthermore, if \( h \) is appropriately chosen, equations (25)–(27) may converge fast. To this end, we have to compute the optimal values of the convergence control parameters from the minimum of the averaged residual errors.

Niu and Chun [37] introduced several methods to obtain the optimal value of \( h. \) The optimal value of the convergence control parameter is defined by using the concept of the square residual error. An error analysis is presented to determine the optimal value of \( h. \) We substitute equations (47)–(49) into equations (4)–(6) and obtain the residual functions as follows:

\[
E_{m,T}(t; h_1) = a \frac{\Delta T}{\Delta t} \psi_T(t; h_1) - p + \mu_T \psi_T(t; h_1) - k_T \psi_T(t; h_1)
\]

\[
+ (1 - \psi_T(t; h_1) + \psi_T(t; h_1)) + k_T \psi_T(t; h_1), \quad (43)
\]

\[
E_{m,I}(t; h_2) = a \frac{\Delta I}{\Delta t} \psi_T(t; h_2) - k_T \psi_T(t; h_2) + \mu_T \psi_T(t; h_2), \quad (44)
\]

\[
E_{m,F}(t; h_3) = a \frac{\Delta F}{\Delta t} \psi_T(t; h_3) - M \psi_T(t; h_3) + k_T \psi_T(t; h_3) + \mu_T \psi_T(t; h_3). \quad (45)
\]

Then, the square residual error for the sixth-order approximation is defined as

\[
SE_{m,T}(h_1) = \frac{1}{(N + 1)} \sum_{l=0}^{N} E_{m,T} \left( \sum_{l=1}^{m} T(l \Delta t) \right), \quad (46)
\]

\[
SE_{m,I}(h_2) = \frac{1}{(N + 1)} \sum_{l=0}^{N} E_{m,I} \left( \sum_{l=1}^{m} I(l \Delta t) \right), \quad (46)
\]

\[
SE_{m,F}(h_3) = \frac{1}{(N + 1)} \sum_{l=0}^{N} E_{m,F} \left( \sum_{l=1}^{m} F(l \Delta t) \right).
\]

The use of the first derivative test enables us to determine the values of the auxiliary parameters \( h_1, h_2, \) and \( h_3 \) for which \( SE_{m,T}(h_1), SE_{m,I}(h_2), \) and \( SE_{m,F}(h_3) \) are minimized. It should be emphasized that the approximation procedures that are used to select the optimal value of \( h \) in FHATM are similar to those of HAM [38].

In Table 3, the minimum values of the square residual error are given for the optimal values of \( h_1, h_2, \) and \( h_3 \) when \( a = \beta = \gamma = 0.99. \)

The absolute residual errors that were calculated for various \( t \in (0, 1) \) are listed in Table 2. These results show that the FHATM obtains an accurate approximate solution for the fractional HIV model (4)–(6). The residual errors are plotted in Figure 4 for \( t \in (0, 1) \) and various values of \( h. \) The square residual errors are plotted in Figure 5, and Figure 6 shows the absolute residual
functions for the optimal $h$. As these figures show, the solution obtained by using FHATM provides us with a sufficiently accurate analytical solution that only requires a few iterative steps. Mathematica software was used to calculate the six-term approximations for $T$, $I$, and $F$, respectively.
Figure 4: Errors of residual functions equations (39)–(41) using the sixth order of the approximation solution for various values of $h$. (a) Comparison errors of residual function $ET$ for various $h$, (b) comparison errors of residual function $EI$ for various $h$, and (c) comparison errors of residual function $EF$ for various $h$.

\[
Y_T(t; h) = \sum_{m=0}^{6} T_m(t) = 1000 + 1.44837 \times 10^{-6}h + 3.70969 \times 10^{-6}h^2 + 5.06867 \times 10^{-6}h^3
\]
\[+ 3.89645 \times 10^{-6}h^4 + 1.59786 \times 10^{-6}h^5 + 2.73079 \times 10^{-7}h^6 + 0.000143992ht^{0.99}
\]
\[+ 0.000377629h^2t^{0.99} + 0.000528248h^3t^{0.99} + 0.000415689h^4t^{0.99} + 0.000174472h^5t^{0.99}
\]
\[+ 0.000305132h^6t^{0.99} + \cdots,
\]

\[
Y_I(t; h) = \sum_{m=0}^{6} I_m(t) = -1.20698 \times 10^{-6}h - 3.0988 \times 10^{-6}h^2 - 4.24412 \times 10^{-6}h^3
\]
\[\[ - 3.27012 \times 10^{-6}h^4 - 1.34403 \times 10^{-6}h^5 - 2.30202 \times 10^{-7}h^6 - 0.000119993ht^{0.99}
\]
\[\[ - 0.000316179h^2t^{0.99} - 0.000351095h^3t^{0.99} - 0.000147925h^4t^{0.99} - 0.000259663h^5t^{0.99} - \cdots,
\]

\[
Y_F(t; h) = \sum_{m=0}^{6} F_m(t) = 0.001 + 0.000146285h + 0.000378272h^2 + 0.000521647h^3
\]
\[\[ + 0.000404615h^4 + 0.000174472h^5t^{0.99} + 0.0388549h^5t^{0.99} + 0.0145343ht^{0.99}
\]
\[\[ + 0.0552967h^3t^{0.99} + 0.0442161h^4t^{0.99} + 0.0188364h^5t^{0.99} + 0.0033401h^6t^{0.99} + \cdots.
\]
Figure 5: Square residual function equations (42)–(44) using the sixth order of the approximation solution for $h \in (−0.9, −0.5)$. (a) Square residual errors of $T$ versus $h$, (b) square residual errors of $I$ versus $h$, and (c) square residual errors of $F$ versus $h$.

Figure 6: Absolute residual function equations (39)–(41) using the sixth order of the approximation solution for $h^*$. (a) Absolute residual error functions for $T$ and the optimal $h$, (b) absolute residual error functions for $I$ and the optimal $h$, and (c) absolute residual error functions for $F$ and the optimal $h$. 
Figure 7: Numerical simulation of the concentrations of uninfected CD4$^+$ T cells $T(t)$, infected CD4$^+$ T cells $I(t)$, and HIV RNA $F(t)$ for different values of $\alpha$, $\beta$, and $\gamma$ and the optimal values of $h^*$. (a) Approximate solutions of $T(t)$, (b) approximate solutions of $I(t)$, and (c) approximate solutions of $F(t)$.

Figure 8: Continued.
Note that, if we set $\alpha = \beta = \gamma = 1$, then the FHATM solution is the same as that obtained with the HAM in [13]. The numerical results are plotted in Figure 7.

5. Numerical Scheme

In this section, we solve fractional a HIV model numerically using the numerical scheme introduced by Toufik and Atangana [27]. The numerical solution and FHATM solution are compared in Figure 8.

6. Conclusion and Further Work

In this study, we successfully solved the fractional HIV infection by using the CD4$^+$ T cells model numerically and analytically, which includes an operator of the type of the Atangana–Bealeau fractional derivative in the Caputo sense (ABC). Analytically approximate solution was obtained for this derivative by incorporating the FHATM in the model of the fractional HIV infection of CD4$^+$ T cells. The solution includes the auxiliary parameter $h$, which provides an easy way to control the convergence region of the resulting infinite series. The results we obtained show that the FHATM is a successful technique for obtaining an approximate solution of the fractional HIV infection of CD4$^+$ T cells. Moreover, our result agrees strongly with the computation of Toufik and Atangana [27]. Studying the dynamics and stability of the system based on the ABC fractional definition and the FHATM algorithm is an interesting idea for the researchers.

Data Availability

No data were used.

Conflicts of Interest

The authors declare that they have no conflicts of interest.

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