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Stability analysis and numerical simulation of fractional model of Leishmaniasis

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Abstract

The disease of leishmaniasis is one that takes some research and study to fully comprehend. As a result, mathematical modeling can be utilized to gain insight into and enhance the accuracy of epidemiological forecasts. A friction model of Leishmaniasis was analyzed using empirical data from Sudan by factoring in the derivatives of Caputo and Atangana-Baleanu. The Caputo and AB derivatives have been subjected to a stability study. A numerical simulation of the suggested ordinary and fractional differential mathematical model follows. The performance was evaluated by calculating its error rating.

Keywords: Mathematical Modelling, Leishmaniosis, Caputo, Atangana–Baleanu derivatives

1. Introduction

Visceral leishmaniasis is a fatal disease transmitted by sandflies. India, Bangladesh, and Nepal have reduced cases of leishmaniasis. Less progress has been made in East Africa, particularly with the continuing epidemic in South Sudan and outbreaks of visceral leishmaniasis. Lack of infrastructure, healthcare personnel, displacement, and malnutrition hamper VL management, diagnostic kits, and medication. However, resistance to pentavalent antimony is a key obstacle that must be overcome before VL may be treated and brought under control. In order to reduce the total amount of time spent

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receiving treatment for this condition, the first line of treatment, which consisted of sodium stibogluconate for 30 days, has been switched out for a more effective injectable combination regimen that includes SSG and PM, which lasts for only 17 days. Relapse in therapy can occur as a consequence of HIV, tuberculosis, malnutrition, or poor treatment, resulting in the parasite remaining in the blood after the first clinical treatment. Malnutrition is another factor that can contribute to relapse in treatment. Because it is difficult to keep track of active patients in Sudan, the country's rates of VL recurrence are unintentionally and passively increasing because the country accepts VL retreat as a part of the total number of VL admissions [1–7].

Over the course of the last three decades, the fractional calculus has emerged as a topic of increasing interest and significance. The fields of physics, chemical engineering, mathematical biology, and even economics all make use of fractional differential equations and nonlinear equations [8–16].

2. Preliminaries

Definition 1. The Riemann-Liouville fractional integral (RLI) operator of order $\alpha > 0$ for a function $y(\tau)$ is given by [17]:

$$
D^{\alpha} y(t) := \frac{1}{\Gamma(n-\alpha)} \int_{0}^{t} (t-\tau)^{n-\alpha-1} y^{n}(\tau) d\tau = I^{n-\alpha} y^{n}(t), t > 0
$$
 (1)

Definition 2. Caputo derivative of order $0 \le n - 1 < \alpha < n$ with the lower limit zero for a function $y(\tau)$ is given by [18]:

$$
I^{\alpha} y(t) := \frac{1}{\Gamma(\alpha)} \int_{0}^{t} (t - \tau)^{\alpha - 1} y(\tau) d\tau, \ t > 0
$$
 (2)

Definition 3. For $y \in H^1(0,t)$, $t > 0$, $T > 0$, $\alpha \in (0,1]$. Then the ABC fractional operator [19] $y(t)$ in the Riemann–Liouville is given by

$$
{}_{0}^{AB}D_{t}^{\alpha}y(t) := \frac{B(\alpha)}{1-\alpha} \frac{d}{dt} \int_{0}^{t} y(\tau) E_{\alpha} \left(\frac{\alpha}{1-\alpha} (t-\tau)^{\alpha} \right) d\tau. \ 0 < \alpha < 1
$$
 (3)

In this expression $B(\alpha)$ satisfies the condition $B(0) = B(1) = 1$.

Definition 4. The Mittag-Leffler function (MLF) is a generalization of the exponential function. This function can be expressed as follows:

$$
E_{\alpha}(t) = \sum_{k=0}^{\infty} \frac{t^k}{\Gamma(\alpha k + 1)}
$$
(4)

3. Anthropological Visceral Leishmaniosis Model with Caputo Derivative

In this Section, the mathematical model of leishmaniasis is a compartmental model with four sub-populations: susceptible, infectious, Recovered, and Recovered and have permanent immunity, for human population and two compartmental for reservoir population Susceptible, infected, in addition to that, we have two compartments for sandflies Susceptible, infected. The human population is the only population in the model that has permanent immunity. The positivity, the number of reproductions, and the equilibrium solutions of the model established in this work have all been determined to be free

of leishmaniasis. Furthermore, the existing cases of leishmaniasis have also been determined along with their respective localities and global stability properties. we obtain the fractional model formulation under Caputo derivative:

$$
s_h(t) = \frac{S_H}{N_H}, i_h(t) = \frac{I_H}{N_H}, p_h(t) = \frac{P_H}{N_H}, r_h(t) = \frac{R_H}{N_H}, s_r(t) = \frac{S_R}{N_R},
$$

$$
I_r(t) = \frac{I_R}{N_R}, s_V(t) = \frac{S_V}{N_V}, i_V(t) = \frac{I_V}{N_V}, s_h(t) = \frac{S_H}{N_H}, m = \frac{N_V}{N_H} \text{ and } N = \frac{N_V}{N_R}
$$

The system of differential equations is given by:

$$
\begin{cases}\n\int_{0}^{C} D_{t}^{\alpha} i_{h} = abmi_{v}N_{h} - \left(\alpha_{1} + \delta + \frac{A_{H}}{N_{H}} - \delta i_{h}\right)i_{h}, \\
\int_{0}^{C} D_{t}^{\alpha} p_{h} = (1 - \sigma)\alpha_{1}i_{h} - \left(\alpha_{2} + \beta + \frac{A_{H}}{N_{H}} - \delta i_{h}\right)p_{h}, \\
\int_{0}^{C} D_{t}^{\alpha} i_{r} = abni_{v}s_{r} - \frac{A_{H}}{N_{H}}i_{r}, \\
\int_{0}^{C} D_{t}^{\alpha} i_{v} = aci_{h}S_{v} + acp_{h}S_{v} + aci_{r}S_{v} - \frac{A_{V}}{N_{V}}i_{v}, \\
\int_{0}^{C} D_{t}^{\alpha} s_{h} = \frac{A_{H}}{N_{H}} - \left[abmi_{v} + \frac{A_{H}}{N_{H}} - \delta i_{h}\right]s_{h}, \\
\int_{0}^{C} D_{t}^{\alpha} r_{h} = \sigma\alpha_{1}i_{h} + (\alpha_{2} + \beta)P_{h} - \left[\frac{A_{H}}{N_{H}} - \delta i_{h}\right]r_{h}, \\
\int_{0}^{C} D_{t}^{\alpha} S_{r} = \frac{A_{R}}{N_{R}} - abni_{v}s_{r} - \frac{A_{H}}{N_{H}}s_{r}, \\
\int_{0}^{C} D_{t}^{\alpha} s_{v} = \frac{A_{V}}{N_{V}} - \left[aci_{h} + acP_{h} + \frac{A_{V}}{N_{V}}\right]s_{v}\n\end{cases}
$$
\n(5)

With initial conditions:

$$
s_h(0) = c_1, i_h(0) = c_2, r_h(0) = c_3, s_r(0) = c_4, I_r(0) = c_5, s_V(0) = c_6, i_V(0) = c_7.
$$

4. Anthropologic Visceral Leishmaniosis Model with ABC Derivative

We obtain the fractional model formulation under Atangana–Baleanu Caputo derivative:

$$
s_h(t) = \frac{S_H}{N_H}, i_h(t) = \frac{I_H}{N_H}, p_h(t) = \frac{P_H}{N_H}, r_h(t) = \frac{R_H}{N_H}, s_r(t) = \frac{S_R}{N_R},
$$

$$
I_r(t) = \frac{I_R}{N_R}, s_V(t) = \frac{S_V}{N_V}, i_V(t) = \frac{I_V}{N_V}, s_h(t) = \frac{S_H}{N_H}, m = \frac{N_V}{N_H} \text{ and } N = \frac{N_V}{N_R}
$$

The system of differential equations is given by:

$$
\begin{cases}\n\frac{AB}{0}D_{t}^{\alpha}i_{h} = abmi_{v}N_{h} - \left(\alpha_{1} + \delta + \frac{A_{H}}{N_{H}} - \delta i_{h}\right)i_{h}, \\
\frac{AB}{0}D_{t}^{\alpha}p_{h} = (1 - \sigma)\alpha_{1}i_{h} - \left(\alpha_{2} + \beta + \frac{A_{H}}{N_{H}} - \delta i_{h}\right)p_{h}, \\
\frac{AB}{0}D_{t}^{\alpha}i_{r} = abni_{v}s_{r} - \frac{A_{H}}{N_{H}}i_{r}, \\
\frac{AB}{0}D_{t}^{\alpha}i_{v} = aci_{h}S_{v} + acp_{h}S_{v} + aci_{r}S_{v} - \frac{A_{V}}{N_{V}}i_{v}, \\
\frac{AB}{0}D_{t}^{\alpha}s_{h} = \frac{A_{H}}{N_{H}} - \left[abmi_{v} + \frac{A_{H}}{N_{H}} - \delta i_{h}\right]s_{h}, \\
\frac{AB}{0}D_{t}^{\alpha}r_{h} = \sigma\alpha_{1}i_{h} + (\alpha_{2} + \beta)P_{h} - \left[\frac{A_{H}}{N_{H}} - \delta i_{h}\right]r_{h}, \\
\frac{AB}{0}D_{t}^{\alpha}S_{r} = \frac{A_{R}}{N_{R}} - abni_{v}s_{r} - \frac{A_{H}}{N_{H}}s_{r}, \\
\frac{AB}{0}D_{t}^{\alpha}s_{v} = \frac{A_{V}}{N_{V}} - \left[aci_{h} + acP_{h} + \frac{A_{V}}{N_{V}}\right]s_{v} \\
\end{cases}
$$
\n(6)

With initial conditions:

$$
s_h(0) = c_1, i_h(0) = c_2, r_h(0) = c_3, s_r(0) = c_4, I_r(0) = c_5, s_V(0) = c_6, i_V(0) = c_7.
$$

5. Stability Analysis

This section covered aspects including the eigenvalues, Jacobian matrix, and equilibrium points of the kala-azar epidemiological model (1).

5.1. Equilibria

5.2. The Jacobian Matrix for the Model

Here, we discuss this epidemiological model stability. The disease free equilibrium point is given as $\mathbf{E}_{1} = (0,0,0,0,1,0,711.58,1)$ and the endemic equilibrium points $\mathbf{E}_{8} = (0,0,0,5.1155e8,0,0,0,0)$.

$$
J(E_1) = \begin{bmatrix}\n-0.031 & 0 & 0 & 0.0157 & 0 & 0 & 0 & 0 \\
0.002 & -0.933 & 0 & 0 & 0 & 0 & 0 & 0 \\
0 & 0 & -4.297e^{-8} & 15907.35 & 0 & 0 & 0 & 0 \\
0.02 & 0.02 & 0.02 & -3.438e^{-11} & 0 & 0 & 0 & 0 \\
0.011 & 0 & 0 & -0.0157 & -4.297e^{-8} & 0 & 0 & 0 \\
0.018 & 0.0933 & 0 & 0 & 0 & -4.297e^{-8} & 0 & 0 \\
0 & 0 & -15907.35 & 0 & 0 & -4.297e^{-8} & 0 \\
-0.02 & -0.02 & 0 & 0 & 0 & 0 & 0 & -3.438e^{-11}\n\end{bmatrix}
$$
\n(7)

5.3. The Basic Reproduction Number

The basic reproduction number is a baseline statistic in epidemiology and is represented by R_0 , which stands for the predicted value of the secondary infections rate per time unit. Using the equation's fractional model (1), We have fours infected classes, rewrite the system of equation (1) for the susceptible and infected classes in the general form:

$$
\frac{dx}{dt} = f(x) - v(x) \tag{9}
$$

Where

$$
f(x) = \begin{pmatrix} abmi_v s_h \\ 0 \\ abmi_v s_r \\ ac(i_h + p_h + i_r) s_v \end{pmatrix}, \text{ and } v(x) = \begin{pmatrix} (\alpha_1 + \delta + \mu_h)i_h \\ (\alpha_2 + \beta + \mu_h) p_h - (1 - \sigma) \alpha_1 i_h \\ \mu_r i_r \\ \mu_v i_v \end{pmatrix}
$$
(10)

Now the Jacobian of $f(x)$ and $v(x)$ of the disease free equilibrium point is:

$$
F = \begin{pmatrix} 0 & 0 & 0 & abm \\ 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & abm \\ ac & ac & ac & 0 \end{pmatrix} \text{ and } V = \begin{pmatrix} \alpha_1 + \delta + \mu_h & 0 & 0 & 0 \\ -(1 - \sigma)\alpha_1 & \alpha_2 + \beta + \mu_h & 0 & 0 \\ 0 & 0 & \mu_r & 0 \\ 0 & 0 & 0 & \mu_v \end{pmatrix}
$$
(11)

we have

$$
R_0 = \rho(FV^{-1}) = \sqrt{\frac{ac[\mu_r abm(\alpha_2 + \delta + \mu_h + (1 - \sigma)\alpha_1) + abn(\alpha_1 + \delta + \mu_h)(\alpha_2 + \delta + \mu_h)]}{\mu_v \ \mu_r(\alpha_1 + \delta + \mu_h)(\alpha_2 + \delta + \mu_h)}}
$$
(12)

Lemma 3.1. The disease-free equilibrium E_0 is locally asymptotically stable if.

 R_0 <1 and unstable if R_0 >1.

6. Simulation

In this section, we simulate our model. Table 1 shows the variable values utilized, and Table 2 shows the parameter values used, together with the initial condition, $s_H(0) = 50$, $P_H(0) = 0$, $i_H(0) = 1$, $r_H(0) = 0$, $s_R(0) = 22$, $I_R(0) = 1$, $s_V(0) = 12$, $i_V(0) = 1$.

Table 2: Parameter values

Parameter	Description	Value	Source
a	Biting rate of sandflies	0.2856 day^{-1}	[16]
$\mathbf b$	Progression rate of VL in sandfly	0.22 day^{-1}	[16]
$\mathbf c$	Progression rate of VL in human and reservoir	0.0714 day^{-1}	[27]
$A_{\rm H}$	Human recruitment rate	10.1009 day^{-1}	Estimated
$A_{\scriptscriptstyle R}$	Reservoir recruitment rate	19.7795 day^{-1}	Estimated
A_{V}	Vector recruitment rate	$38858.62 \text{ day}^{-1}$	Estimated
μ_h	Natural mortality rate of humans	$4.341e-6$ day ⁻¹	$[2]$
μ_r	Natural mortality rate of reservoirs	0.0017 day^{-1}	$[1]$
μ_{v}	Natural mortality rate of vectors	0.0668 day^{-1}	$[1]$
α_{1}	Treatment rate of VL	0.02	$[2]$
α_{2}	PKDL recovery rate without treatment	0.033	[42]
σ	Recovery rate from VL infection after treatment	0.9	$[1] % \includegraphics[width=0.9\columnwidth]{figures/fig_10.pdf} \caption{The figure shows the number of times of the estimators in the left and right.} \label{fig:time} %$
$1-\sigma$	Developing PKDL rate after treatment	0.1	$[1]$
δ	Death rate due to VL	0.011	$[35]$
β	PKDL recovery rate after treatment	0.9	$[1] % \includegraphics[width=0.9\columnwidth]{figures/fig_10.pdf} \caption{The figure shows the number of times of the estimators in the left and right.} \label{fig:time} %$

λ_i	Eigenvalues	Stability
λ_1	$(17.833, 0, 0, 0, 0, 0, 0, -17.833)$	Unstable
λ_2	$\left(-3.438e^{-11},-2.8e^{-8},-2.8e^{-8},-4.297e^{-8},-0.31,-0.933,-035456.6,-1.14e^{10}\right)$	Stable
	2.2 50 ГC $-c$ 49.5 $- ABC$ $-$ ABC $\sqrt{2}$ 49 $\begin{array}{c}\n48.5 \\ 48.5 \\ 447 \\ 47 \\ 50 \\ 60 \\ 60 \\ 70 \\ 80 \\ 80 \\ 80 \\ 90 \\ 80 \\ $ R and R 1.6 R 1.6 R 1.4 46.5 46 $1.2\,$ 45.5 $45\genfrac{}{}{0pt}{}{\rangle}{0}$ $\mathbf{1}$ 0.2 0.3 $\mathbf 0$ 0.1 0.4 $0.5\,$ 0.6 0.7 0.8 0.9 0.1 0.2 0.3 0.4 $0.5\,$ 0.6 0.7 0.8 0.9 $\mathbf{1}$ \mathfrak{t} $^\mathrm{t}$ 10 $-c$ $= ABC$ $-c$ $- ABC$ $\boldsymbol{9}$ Recovered and have permanent immunity $\begin{array}{ccc}\n0 & 2 \\ 0 & 3\n\end{array}$ $\begin{array}{ccc}\n0 & 2 \\ 0 & 3\n\end{array}$ 8 $\overline{}$ Infected humans $\,$ 6 $\,$ 5 $\overline{4}$ $\sqrt{3}$	$\mathbf{1}$
	\overline{c} $0.2\frac{L}{0}$ $\mathbf{1}$ 0.5 $\mathbf{0}$ 0.2 0.3 0.4 0.6 $0.8\,$ 0.9 0.1 0.2 $0.3\,$ 0.4 0.5 $0.6\,$ 0.7 $0.8\,$ 0.9 0.1 $0.7\,$ 1 \mathfrak{t} 22 4 $\frac{C}{E}$ ∎с $- ABC$ 3.5 21.5 Susceptible reservoir 20 20 20 $\sqrt{3}$ Infected reservoir 2.5 ᄾ $\overline{\mathbf{c}}$ 19.5 $1.5\,$ 19 0.3 0.6 0.7 0.8 $\overline{0}$ 0.2 0.3 $0.5\,$ 0.6 0.7 0.8 0.1 0.2 0.4 0.5 0.9 $\mathbf{1}$ 0.1 0.4 0.9 $\mathbf{0}$ \mathfrak{t} $^\mathrm{t}$	1 $\mathbf{1}$
	9 12 -c ĿС $- ABC$ $- ABC$ 11.8 $\bf8$ 11.6 $\overline{\mathfrak{c}}$ se 11.4 The Samples 11.2 The 11.2 The 10.8 The 10.8 Infected sandflies $\,$ 6 5 $\sqrt{4}$ $\sqrt{3}$ 10.6 $\sqrt{2}$ 10.4 $10.2 - 0$ $\mathbf{1}$ 0.4 0.5 $0.6\,$ 0.7 $0.8\,$ 0.9 $0.6\,$ 0.8 $\mathbf 0$ 0.2 0.3 0.1 0.2 0.3 0.4 $0.5\,$ 0.7 0.9 0.1 1	1

Table 4: The eigenvalues corresponding to matrix *J* are.

Figure 1: Figures of the systems of fractional orders model for $\alpha = 0.99$.

Figure 2: Figures of the fractional orders model for $\alpha = 1$.

7. Conclusion

We used MATLAB to simulate the Caputo derivative and the AB derivative-based fractional model. Furthermore, the model calculations and the corresponding graphs of the fractional derivative provide a comprehensive explanation of Leishmaniosis. The use of fractional derivatives is suggested for a more accurate depiction of the Leishmaniosis pandemic. The AB derivative's superior predictive power may be traced back to the fact that the relevant model has a non-singular kernel.

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