Optimal Control and Sensitivity Analysis of a Fractional Order TB Model

Silvério Rosa¹, Delfim F. M. Torres ²,*

¹Department of Mathematics and Instituto de Telecomunicações (IT), University of Beira Interior, 6201-001 Covilhã, Portugal
²Center for Research and Development in Mathematics and Applications (CIDMA), Department of Mathematics, University of Aveiro, 3810-193 Aveiro, Portugal

Abstract A Caputo fractional-order mathematical model for the transmission dynamics of tuberculosis (TB) was recently proposed in [Math. Model. Nat. Phenom. 13 (2018), no. 1, Art. 9]. Here, a sensitivity analysis of that model is done, showing the importance of accuracy of parameter values. A fractional optimal control (FOC) problem is then formulated and solved, with the rate of treatment as the control variable. Finally, a cost-effectiveness analysis is performed to assess the cost and the effectiveness of the control measures during the intervention, showing in which conditions FOC is useful with respect to classical (integer-order) optimal control.

Keywords Tuberculosis; Compartmental mathematical models; Fractional optimal control.

AMS 2010 subject classifications 34A08, 49M05, 92C60

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1. Introduction

Tuberculosis (TB) is an infectious disease caused by the bacterium Mycobacterium tuberculosis, which is usually spread through the air when people who have active TB in their lungs cough, spit, speak, or sneeze. TB is one of the top ten causes of death worldwide, which justifies the interest of researchers on the area: see, e.g., [17, 19, 23]. A good survey on optimal models of TB is presented in [20]. Delays are introduced in a TB model in [17], representing the time delay on the diagnosis and initiation of treatment of individuals with active TB infection. Optimal control strategies to minimize the cost of interventions, considering reinfection and post-exposure interventions, are investigated in [19]. In [15], the potential of two post-exposure interventions, treatment of early latent TB individuals and prophylactic treatment/vaccination of persistent latent TB individuals, is investigated. A mathematical model for TB is studied in [5] from the optimal control point of view, using a multiobjective optimization approach. For numerical simulations using TB real data from Angola, see [18]. In spite of the numerous works on tuberculosis, the literature on fractional-order mathematical models for TB is still scarce. In [21], it is proposed a multi-strain TB model with variable-order fractional derivatives and a numerical scheme to approximate the endemic solution, numerically, is developed. More recently, in 2018, a Caputo type fractional-order mathematical model for the transmission dynamics of tuberculosis was proposed, based on the nonlinear differential system studied in [25], and its stability is investigated [24]. Such model is here subject to fractional optimal control theory [4, 7], and its usefulness to tackle a TB epidemic scenario discussed.

The paper is organized as follows. In Section 2, we introduce the fractional-order TB model. The main results are then given in Section 3: sensitivity analysis of the fractional TB model (Section 3.1); fractional optimal control,
cost-effectiveness and numerical simulations for the TB model (Sections 3.2 and 3.3). We end with Section 4 of conclusions.

2. Fractional-order Tuberculosis Model

In this section, we consider a Caputo fractional-order tuberculosis (TB) model due to [24]. The model describes the dynamics of a population that is susceptible to infection by the *Mycobacterium tuberculosis* with incomplete treatment. The population consists of four compartments: susceptible individuals (S); latent individuals (L), which have been infected but are not infectious and do not show symptoms of the disease; infectious individuals (I), which have active TB, may transmit the infection, but are not under treatment; and infected individuals in treatment (T).

The susceptible population is increased by the recruitment of individuals into the population at a rate $\Lambda$. All individuals are exposed to natural death, at a constant rate $\mu$. Deterministic continuous transitions between the compartments, also known as states, are used. Susceptible individuals acquire TB infection by the contact with infected individuals at a rate $\beta I$, where parameter $\beta$ is the transmission coefficient. Individuals in the latent class, $L$, become infectious at a rate $\varepsilon$, and infectious individuals, $I$, start treatment at a rate $\gamma$. Treated individuals, $T$, leave their compartment at rate $\delta$. After leaving the treatment compartment, an individual may enter compartment $L$, due to the remainder of *Mycobacterium tuberculosis*, or compartment $I$, due to the failure of treatment. The parameter $k$, $0 \leq k \leq 1$, represents the failure of the treatment, where $k = 0$ means that all the treated individuals shall become latent, while $k = 1$ means that the treatment fails and all the treated individuals shall still be infectious. Infectious, $I$, and under treatment individuals, $T$, may suffer TB-induced death at the rates $\alpha_1$ and $\alpha_2$, respectively.

The Caputo fractional-order system of differential equations that describes the above assumptions is

\[
\begin{align*}
\frac{\mathcal{C}_0^\alpha D_t^\alpha S(t)}{S(t)} &= \Lambda - \beta I(t)S(t) - \mu S(t), \\
\frac{\mathcal{C}_0^\alpha D_t^\alpha L(t)}{L(t)} &= \beta I(t)S(t) + (1-k)\delta T(t) - (\mu + \varepsilon)L(t), \\
\frac{\mathcal{C}_0^\alpha D_t^\alpha I(t)}{I(t)} &= \varepsilon L(t) + k\delta T(t) - (\mu + \gamma + \alpha_1)I(t), \\
\frac{\mathcal{C}_0^\alpha D_t^\alpha T(t)}{T(t)} &= \gamma I(t) - (\mu + \delta + \alpha_2)T(t),
\end{align*}
\]

where $\frac{\mathcal{C}_0^\alpha D_t^\alpha}{\alpha}$ denotes the left Caputo derivative of order $\alpha \in (0, 1)$ [12, 16]. Note that when $\alpha = 1$, the fractional compartmental model (1) represents the classical TB model studied in [25]. In Figure 1, one can find a diagram representing the model dynamics.

![Figure 1. Epidemiological scheme of the mathematical model (1).](image-url)
3. Main results

We begin by investigating the sensitiveness of the TB model (1), discussed in Section 2, with respect to the variation of each one of its parameters.

3.1. Sensitivity analysis

One of the most significant thresholds when studying infectious disease models is the basic reproduction number \( R_0 \). The basic reproduction number for the TB model (1) is given by

\[
R_0 = \frac{\beta \varepsilon b_3 \Lambda}{\mu b_1 b_2 b_3 - \mu \delta \gamma (1 - k) \varepsilon + kb_2} \tag{2}
\]

with \( b_1 = \mu + \varepsilon, b_2 = \mu + \gamma + \alpha_1 \) and \( b_3 = \mu + \delta + \alpha_2 \) [25].

Now we perform a sensitivity analysis for the endemic threshold (2). Such analysis tells us how important each parameter is to disease transmission. This information is crucial not only for experimental design, but also to data assimilation and reduction of complex models [13]. Sensitivity analysis is commonly used to determine the robustness of model predictions to parameter values, since there are usually errors in collected data and presumed parameter values. It is used to discover parameters that have a high impact on the threshold \( R_0 \) and should be targeted by intervention strategies. More accurately, sensitivity indices’s allows us to measure the relative change in a variable when parameter changes. For that purpose we use the normalized forward sensitivity index of a variable, with respect to a given parameter, which is defined as the ratio of the relative change in the variable to the relative change in the parameter. If such variable is differentiable with respect to the parameter, then the sensitivity index is defined using partial derivatives.

**Definition 1 (See [3, 14])**

The normalized forward sensitivity index of \( R_0 \), which is differentiable with respect to a given parameter \( p \), is defined by

\[
\Upsilon_{R_0}^p = \frac{\partial R_0}{\partial p} \frac{p}{R_0}.
\]

The values of the sensitivity indices for the parameters values of Table 1, are presented in Table 2.

Table 1. Values of the models’ parameters taken from [24, 25].

<table>
<thead>
<tr>
<th>Name</th>
<th>Description</th>
<th>Value</th>
</tr>
</thead>
<tbody>
<tr>
<td>( \Lambda )</td>
<td>Recruitment rate</td>
<td>792.8571</td>
</tr>
<tr>
<td>( \beta )</td>
<td>Transmission coefficient</td>
<td>0.0005</td>
</tr>
<tr>
<td>( \mu )</td>
<td>Natural death rate</td>
<td>1/70</td>
</tr>
<tr>
<td>( k )</td>
<td>Treatment failure rate</td>
<td>0.15</td>
</tr>
<tr>
<td>( \delta )</td>
<td>Rate at which treated individuals leave the ( T ) compartment</td>
<td>1.5</td>
</tr>
<tr>
<td>( \varepsilon )</td>
<td>Rate at which latent individuals ( L ) become infectious</td>
<td>0.00368</td>
</tr>
<tr>
<td>( \alpha_1 )</td>
<td>TB-induced death rate for infectious individuals ( I )</td>
<td>0.3</td>
</tr>
<tr>
<td>( \alpha_2 )</td>
<td>TB-induced death rate for under treatment individuals ( T )</td>
<td>0.05</td>
</tr>
</tbody>
</table>

Note that the sensitivity index may depend on several parameters of the system, but also can be constant, independent of any parameter. For example, \( \Upsilon_{\Lambda}^{R_0} = +1 \), meaning that increasing (decreasing) \( \Lambda \) by a given percentage increases (decreases) always \( R_0 \) by that same percentage. The estimation of a sensitive parameter should be carefully done, since a small perturbation in such parameter leads to relevant quantitative changes. On the other hand, the estimation of a parameter with a rather small value for the sensitivity index does not require as much attention to estimate, because a small perturbation in that parameter leads to small changes [10]. According to our results in Table 2, special attention should be paid to the estimation of the death rate, \( \mu \). In contrast, the rate at which individuals leave state \( T, \delta \), does not require as much attention because of its low value of the sensitivity.
Table 2. Sensitivity of $R_0$ evaluated for the parameter values given in Table 1.

<table>
<thead>
<tr>
<th>Parameter</th>
<th>Sensitivity index</th>
</tr>
</thead>
<tbody>
<tr>
<td>$\mu$</td>
<td>−1.93223</td>
</tr>
<tr>
<td>$\varepsilon$</td>
<td>+0.911803</td>
</tr>
<tr>
<td>$\gamma$</td>
<td>−0.605532</td>
</tr>
<tr>
<td>$\alpha_1$</td>
<td>−0.376538</td>
</tr>
<tr>
<td>$\delta$</td>
<td>0.0112215</td>
</tr>
<tr>
<td>$\alpha_2$</td>
<td>−0.0872783</td>
</tr>
<tr>
<td>$\Lambda$</td>
<td>+1</td>
</tr>
<tr>
<td>$k$</td>
<td>+0.100487</td>
</tr>
<tr>
<td>$\beta$</td>
<td>+1</td>
</tr>
</tbody>
</table>

index. This is well illustrated in Figure 2, where we can see in Figure 2a the graphics of the number of infective with and without an increment of 15% for the parameter $\mu$ (most sensitive parameter), and in Figure 2b with identical comparison of graphics for parameter $\delta$ (less sensitive parameter).

Using the values of the parameters as proposed in [24], we obtain results very different from those that the authors present there. We noticed that the value of parameter $\mu$ presented in [24] ($\mu' = 0.143$), the most sensitive one, is erroneous, being 10 times the correct value [25] and was not the value that authors’ used in the simulation. This particular case highlights the importance of the analysis we have carried out here.

3.2. Fractional optimal control

The circumstances on which variation of the variables of the model depend, may be controlled. In case of TB disease, the fraction of infectious individuals that is identified and put under treatment is one of the most commonly used. Therefore, the parameter $\gamma$ is, in what follows, replaced by a control variable. Consequently, we consider the following fractional optimal control problem: to minimize the number of infectious individuals and the cost of the control of the disease with the identification and treatment of the patients, that is,

$$
\min \mathcal{J}(I(t), u(t)) = \int_0^T I(t) + B\rho u(t)^2 \, dt
$$

(3)
with $0 < B < \infty$, subject to the fractional control system

\[
\begin{align*}
\overset{\alpha}{D}t S(t) & = \Lambda - \beta I(t)S(t) - \mu S(t) \\
\overset{\alpha}{D}t L(t) & = \beta I(t)S(t) + (1 - k)\delta T(t) - (\mu + \varepsilon)L(t) \\
\overset{\alpha}{D}t I(t) & = \varepsilon L(t) + k\delta T(t) - (\mu + u(t) + \alpha_1)I(t) \\
\overset{\alpha}{D}t T(t) & = u(t)I(t) - (\mu + \delta + \alpha_2)T(t)
\end{align*}
\]

and given initial conditions

\[S(0), L(0), I(0), T(0) \geq 0.\]  

Here, $\rho$ is the maximum number of infectious individuals of the problem without control; and the control variable, $u$, designate the fraction of individuals that is put under treatment. The set of admissible control functions is

\[\Omega = \{u(\cdot) \in L^\infty(0, t_f) : 0 \leq u(t) \leq u_{\max}, \forall t \in [0, t_f]\}.\]

Pontryagin’s maximum principle (PMP) for fractional optimal control can be used to solve the problem [1, 2, 8, 9]. The Hamiltonian associated with our optimal control problem is

\[H = I + B \rho u^2 + p_1(\Lambda - \beta IS - \mu S) + p_2(\beta IS + (1 - k)\delta T - (\mu + \varepsilon)L) + p_3(\varepsilon L + k\delta T - (\mu + u + \alpha_1)I) + p_4(uI - (\mu + \delta + \alpha_2)T);\]

the adjoint system uphold that the co-state variables $p_i(t), i = 1, \ldots, 4$, verify

\[
\begin{align*}
\overset{\alpha}{D}t p_1(t) & = \mu p_1(t) - \beta I(t)p_1(t) - p_2(t), \\
\overset{\alpha}{D}t p_2(t) & = (\varepsilon + \mu)p_2(t) - \varepsilon p_3(t), \\
\overset{\alpha}{D}t p_3(t) & = -1 + (\alpha_1 + u(t) + \mu)p_3(t) - u(t)p_4(t) + \beta S(t)(p_1(t) - p_2(t)), \\
\overset{\alpha}{D}t p_4(t) & = (\alpha_2 + \mu + \delta)p_4(t) + \delta(k - 1)p_2(t) - \delta k p_3(t),
\end{align*}
\]

which is a fractional system of right Riemann–Liouville fractional derivatives, $\overset{\alpha}{D}t p_i$. In turn, the optimality condition of PMP establishes that the optimal control is given by

\[u(t) = \min \left\{ \max \left\{ 0, \frac{(p_3(t) - p_4(t))I(t)}{2B\rho} \right\}, u_{\max} \right\}.\]

In addition, the following transversality conditions hold:

\[\overset{\alpha}{D}t^{1-\alpha} p_i \big|_{t_f} = 0 \iff I^{1-\alpha}_t p_i \big|_{t_f} = 0, \quad i = 1, \ldots, 4,\]

where $I^{1-\alpha}_t$ is the right Riemann–Liouville fractional integral of order $1 - \alpha$.

### 3.3. Numerical results and cost-effectiveness of the fractional TB optimal control problem

The PMP is used to numerically solve the optimal control problem (3)–(6), as discussed in Section 3.2, in the classical ($\alpha = 1$) and fractional ($0 < \alpha < 1$) cases, using the predict-evaluate-correct-evaluate (PECE) method of Adams–Basforth–Moulton [6], implemented in MATLAB. First, we solve system (4) by the PECE procedure with initial values for the state variables given by Table 3 and a guess for the control over the time interval $[0, t_f]$, and obtain the values of the state variables $S, L, I$ and $T$. A change of variables is applied to the adjoint system (7) and to the transversality conditions (9), obtaining a left Riemann–Liouville fractional initial value problem. Such system is solved with the PECE procedure and the values of the co-state variables $p_i, i = 1, \ldots, 4$, are obtained. The control is then updated by a convex combination of the previous control and the value computed according to (8). This procedure is repeated iteratively until the values of the controls at the previous iteration are very close to...
the ones at the current iteration. The solutions of the classical problem \((\alpha = 1)\) where successfully confirmed by an algorithm that uses a classical forward-backward scheme, also implemented in MATLAB.

In our numerical experiments, we consider that the maximum number of infectious individuals for the problem without control, \(\rho\), is 452.758, \(u_{\text{max}} = 1\), \(B = 0.15\), while the other parameters are fixed according with Table 1. Note that, with the above values for the parameters, \(R_0 > 1\) (endemic situation). Our initial conditions, given by Table 3, guarantee the existence of a non-trivial endemic equilibrium for system (4) without control. Because the WHO (World Health Organization) goals for most diseases are usually fixed for five years periods, we considered \(t_f = 5\).

Table 3. Initial conditions for the fractional optimal control problem (3)–(6) with parameters given by Table 1, corresponding to the endemic equilibrium of TB model (1).

<table>
<thead>
<tr>
<th>(S(0))</th>
<th>(L(0))</th>
<th>(I(0))</th>
<th>(T(0))</th>
</tr>
</thead>
<tbody>
<tr>
<td>7779.28</td>
<td>43511.9</td>
<td>175.267</td>
<td>78.4299</td>
</tr>
</tbody>
</table>

Without loss of generality, we considered the fractional order derivatives \(\alpha = 1.0, 0.9\) and 0.8. In Figures 3 and 4a, we find the solutions of the fractional optimal control problem for those values of \(\alpha\). We can see that a change in the value of \(\alpha\), corresponds significant variations of the state and control variables. The existence of an endemic situation \((R_0 > 1)\), a very large initial number of latent individuals (about 84% of population), and a control that vanishes at the end of the time interval motivates that, in the end of the time interval, the number of infected individuals exceeds its initial value. Beyond those values of \(\alpha\), others values less than one were also tested, but the results do not change qualitatively. We note that, in all the experiments, decreasing \(\alpha\) meant increasing the number of infected, as Figure 3c evidences.

In Figure 4b, the efficacy function [15] is displayed. It is defined by

\[
F(t) = \frac{I(0) - I^*(t)}{I(0)} = 1 - \frac{I^*(t)}{I(0)},
\]

where \(I^*(t)\) is the optimal solution associated with the fractional optimal control and \(I(0)\) is the corresponding initial condition. This function measures the proportional variation in the number of infectious individuals after the application of the control \(u^*\), by comparing the number of infected individuals at time \(t\) with the initial value \(I(0)\). We observe that \(F(t)\) exhibits the inverse tendency of \(I(t)\). Once \(I(t)\) ends with values bigger than its initial value, \(F(t)\) ends with negative values.

To assess the cost and the effectiveness of the proposed fractional control measure during the intervention period, some summary measures are presented. The total cases averted by the intervention, during the time period \(t_f\), is defined in [15] by

\[
A = t_f I(0) - \int_0^{t_f} I^*(t) \, dt,
\]

where \(I^*(t)\) is the optimal solution associated with the fractional optimal control \(u^*\) and \(I(0)\) is the corresponding initial condition. Note that this initial condition is obtained as the equilibrium proportion \(\bar{T}\) of system (4) without treatment intervention, which is independent on time, so that \(t_f I(0) = \int_0^{t_f} \bar{T} \, dt\) represents the total infectious cases over the given period of \(t_f\) years. We define effectiveness as the proportion of cases averted on the total cases possible under no intervention [15]:

\[
\bar{F} = 1 - \frac{\int_0^{t_f} I^*(t) \, dt}{t_f I(0)}.
\]

The total cost associated with the intervention is defined in [15] by

\[
TC = \int_0^{t_f} C \, u^*(t) I^*(t) \, dt,
\]

Figure 3. State variables of the fractional optimal control problem (3)–(6), with values from Table 1, weight $B = 0.15$, and the fractional order derivatives $\alpha = 1.0, 0.9$ and $0.8$.

Figure 4. Control function $u(t)$ and efficacy function $F(t)$ associated to the fractional optimal control problem (3)–(6), with values from Table 1, weight $B = 0.15$, and the fractional order derivatives $\alpha = 1.0, 0.9$ and $0.8$.

where $C$ corresponds to the unit cost, per person, of detection and treatment of infectious individuals. Following [11, 15], the average cost-effectiveness ratio is given by

$$ACER = \frac{TC}{A}.$$
In Table 4, the cost-effectiveness measures, for the fractional optimal control problem we have analysed, are summarized. These results show the effectiveness of the control to reduce TB infectious individuals and the superiority of the classical model ($\alpha = 1$).

Table 4. Summary of cost-effectiveness measures for classical and fractional ($0 < \alpha < 1$) TB disease optimal control problems. Parameters according to Tables 1 and 3 with $C = 1$.

<table>
<thead>
<tr>
<th>$\alpha$</th>
<th>$A$</th>
<th>$TC$</th>
<th>$ACER$</th>
<th>$\overline{F}$</th>
</tr>
</thead>
<tbody>
<tr>
<td>1.0</td>
<td>88.6409</td>
<td>617.147</td>
<td>6.96233</td>
<td>0.101150</td>
</tr>
<tr>
<td>0.9</td>
<td>73.4419</td>
<td>608.801</td>
<td>8.28956</td>
<td>0.083806</td>
</tr>
<tr>
<td>0.80</td>
<td>54.8307</td>
<td>597.161</td>
<td>10.891</td>
<td>0.062568</td>
</tr>
</tbody>
</table>

The impact of the variation of the weight $B$ on the cost functional is displayed in Figure 5a, and the effectiveness measure $\overline{F}$ is displayed in Figure 5b. When the cost of treatment increases, we observe that the fractional model can be more effective in reducing the number of infective individuals at a not so high cost (Figure 5a).

![Figure 5a](image1.png)  ![Figure 5b](image2.png)

(a) Evolution of the cost functional $J$.  
(b) Evolution of the effectiveness $\overline{F}$.

Figure 5. Impact of the variation of the weight $B$ on the cost functional value, $J$, (left) and on the effectiveness measure $\overline{F}$ (right) for fractional order derivatives $\alpha = 1.0$, 0.9 and 0.8.

4. Conclusions

A sensitivity analysis was carried out for the basic reproduction number of the TB model [24], proving that the most important parameter to have into account is the natural death rate $\mu$. The application of optimal control to the fractional TB model shows that treatment is effective on the reduction of infected individuals. A change in the value of the fractional derivative order, $\alpha$, corresponds to substantial variations on the solutions of the fractional optimal control problem. The fractional model ($0 < \alpha < 1$) is recommended when treatment is expensive, and only in that case, because in that situation it is more effective and not so costly as the classical model.

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