

### **Research** Article

## A Model for the Propagation and Control of Pulmonary Tuberculosis Disease in Kenya

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Pulmonary tuberculosis is among the leading infectious diseases causing mortality worldwide. Therefore, scaling up intervention strategies to reduce the spread of infections in the population is imperative. In this paper, a population-based compartmental approach has been employed to formulate a mathematical model of pulmonary tuberculosis that incorporates an asymptomatic infectious population. The model includes asymptomatic infectious individuals since they spread infections incessantly to susceptible populations without being noticed, thus contributing to the high rate of infection transmission. Qualitative and numerical analyses were performed to determine the impact of various intervention strategies on controlling infection transmission in the population. Sensitivity and numerical results indicate that increasing screening of latently infected and asymptomatic infectious individuals reduces infection transmission to the susceptible population. Numerical results demonstrate that the combination of vaccination, screening, and treatment of all forms of pulmonary tuberculosis is the most effective intervention in decreasing infection transmission. Furthermore, a combination of screening and treatment of all forms of pulmonary tuberculosis proves more effective than a combination of vaccination and treatment of symptomatic infectious individuals alone. Treating the symptomatic infectious population. These study findings will guide healthcare officials in making decisions regarding the screening of latently infected and asymptomatic infectious pulmonary tuberculosis patients, thereby aiding in the fight against epidemics of this disease.

#### 1. Introduction

Mathematical modeling is one of the valuable tools that explore the transmission dynamics of infectious diseases and assess the impact of various control interventions. Therefore, mathematical modeling of infectious diseases is used to guide public health policies and inform decision-making during epidemics [1, 2]. This paper employs a classical SEIR transmission mechanism to formulate a novel model for pulmonary tuberculosis, aiming to accurately depict its natural progression. The SEIR model, a widely used compartmental model in epidemiology, is utilized to forecast the spread of infectious diseases with latent or exposed phases [3]. During the latent phase of a disease, individuals are neither infectious nor symptomatic [4]. The evolution of infectious diseases, taking into consideration the SEIR model, has been discussed by several authors [5–9]. Research suggests that around one-quarter of the global population harbors latent pulmonary tuberculosis infections [10]. Consequently, an epidemiological model for pulmonary tuberculosis must account for this latent population and hence adopts the classical SEIR framework. Within the SEIR model, the population is categorized into four compartments: Susceptible (S), Exposed (E), Infectious (I), and Recovered (R) [2].

Pulmonary tuberculosis is a contagious disease that primarily affects the lungs. It is caused by microorganisms known as Mycobacterium tuberculosis, which spread when an infectious individual coughs, sneezes, speaks, or sings [4, 10]. The disease exerts a heavy toll on the human population, causing a significant number of deaths worldwide. Every year, approximately 10,000,000 people fall ill with pulmonary tuberculosis across the globe, with about 15% of them succumbing to the disease. In 2022, approximately 10,600,000 people fell ill with tuberculosis worldwide, resulting in 1.6 million deaths [11]. In Kenya, data from Civil Registration Services show that pulmonary tuberculosis accounted for about 5.4% of all reported deaths in 2019 [12]. Therefore, it is imperative to scale up preventive measures aimed at reducing the transmission of infections to the susceptible population. The United Nations' "Agenda 2030 Sustainable Development Goals (SDGs)" has identified tuberculosis as one of the communicable diseases that need to be eradicated worldwide by 2030 for sustainable development [13].

Various mathematical models for pulmonary tuberculosis disease have been formulated, focusing on diverse strategies to control the transmission of infections, such as vaccination, early diagnosis, social protection, and treatment of drug-resistant strains [14-16]. Athithan et al. [17] analyzed a nonlinear model of pulmonary tuberculosis considering case detention and treatment, and their results showed that sustaining treatment and case detention hold great promise in controlling the spread of infections. Salpeter et al. [18] conducted a study on the mathematical modeling of pulmonary tuberculosis with estimates of reproduction number and infection delay function. Their outcome revealed that the risk of infection reactivation decreases rapidly and then gradually, for the first ten years after infection. Houben et al. [19] formulated a mathematical model to estimate the global burden of latent tuberculosis infection, and their results estimated that approximately a quarter of the world's population was infected with latent tuberculosis in 2014. Vaccination and effective contact rates on the spread of pulmonary tuberculosis were assessed using a mathematical model in [20, 21]. Their results showed that vaccination coverage is not sufficient to control pulmonary tuberculosis, and the effective contact rate has a higher impact on the spread of infections. Aparicio et al. [22] explored the strengths and limitations of homogeneous and heterogeneous mixing in tuberculosis epidemics through mathematical modeling, and their results indicated that a decrease in pulmonary tuberculosis incidence was due to a reduction in progression rates. Kasereka et al. [23] simulated a mathematical model of pulmonary tuberculosis transmission in the Democratic Republic of Congo, revealing that monitoring contacts, detection of latent infection, and treatment are the optimal strategies to reduce the transmission of infections in the population.

The main purpose of this study is to investigate the impact of various intervention strategies on controlling the transmission of infections in the population. We formulated and analyzed a deterministic pulmonary tuberculosis model incorporating an asymptomatic infectious population. The justification for incorporating an asymptomatic infectious population stems from a report by the National TB Prevalence Survey of 2016 in Kenya, which showed that 26% of prevalent cases diagnosed during their survey were asymptomatic infectious [24]. We investigated the effects of screening asymptomatic and latently infected populations on controlling the transmission of infections to the susceptible population. This paper is organized as follows: Section 2 presents the model formulation, while in Section 3 mathematical analyses have been performed. Section 4 provides the numerical simulations of the model to illustrate the impact of various intervention strategies on controlling the transmission of infections. Finally, Section 5 presents the conclusion.

#### 2. Pulmonary Tuberculosis Model Formulation Incorporating Asymptomatic Infectious Population

This study adopts a classical SEIR transmission mechanism to formulate a new epidemiological model describing the transmission of pulmonary tuberculosis disease. The model incorporates an asymptomatic infectious population, as they spread infections incessantly to the susceptible population without being noticed, thus contributing to the high rate of transmission. The total human population size at a time t, denoted as N(t), is subdivided into susceptible S(t), vaccinated V(t), latent infected E(t), asymptomatic infectious  $I_a(t)$ , symptomatic infectious  $I_s(t)$ , latent infected undergoing treatment  $T_a(t)$ , symptomatic infectious undergoing treatment  $T_s(t)$ , and recovered R(t). Hence, for total human population, we have N(t) = S(t) + V(t) + E(t) $+ I_a(t) + I_s(t) + T_E(t) + T_a(t) + T_s(t) + R(t)$ .

Individuals are recruited into the population at a constant rate of  $\pi$ . A fraction of the recruits is vaccinated at a constant rate of P and enters the vaccinated class, whereas the rest become susceptible. The susceptible individuals become latently infected after effective contact with any of the following populations: symptomatic infectious, asymptomatic infectious, symptomatic infectious undergoing treatment, and asymptomatic infectious undergoing treatment. The force of infection is represented by  $\lambda = \beta (I_s + \eta_1 I_a + \eta_2 T_s + \eta_3 T_a)$ , where  $\beta$  represents the transmission rate of pulmonary tuberculosis infections, while  $\eta_1$ ,  $\eta_2$ , and  $\eta_3$  are the dimensionless transmission coefficients accounting for the relative infectiousness of asymptomatic infectious individuals, symptomatic infectious individuals undergoing treatment, and asymptomatic infectious individuals undergoing treatment, respectively, with  $\eta_3 < \eta_2 < \eta_1$ . This hierarchy assumes that asymptomatic infectious individuals  $(I_a)$  are more infectious than symptomatic individuals undergoing treatment  $(T_s)$ , who are, in turn, more infectious than asymptomatic infectious individuals undergoing treatment  $(T_a)$ . The dimensionless transmission coefficients,  $\eta_1$ ,  $\eta_2$ , and  $\eta_3$  are considered to be less than 1. The model assumes that vaccination is not 100% effective, and thus the vaccinated class has a chance of being latently infected at a force of infection given by  $\lambda_{\nu} = (1 - \rho)\lambda = (1 - \rho)\beta$  $(I_s + \eta_1 I_a + \eta_2 T_s + \eta_3 T_a)$ , where  $\rho$  is the vaccine efficacy, such that  $0 \le \rho \le 1$ . The latent infected individuals, denoted by E(t), are either screened at a constant rate of  $\theta_1$  and moved to the latent infected undergoing treatment class  $T_E(t)$  or progress to the asymptomatic infectious class  $I_a(t)$ , at a rate of  $\chi_1$ . Alternatively, they may develop severe disease and transition to the symptomatic infectious class  $I_s(t)$ , at a constant rate of  $\chi_2$ . Asymptomatic infectious individuals are either screened at a rate of  $\theta_2$  and moved to the asymptomatic infectious undergoing treatment class  $T_a(t)$ , or they progress to severe disease and join the symptomatic infectious class  $I_{s}(t)$  at a rate of  $\omega$ . Symptomatic infectious individuals are identified for treatment at a rate of  $\theta_3$  and moved to the symptomatic infectious undergoing treatment class  $T_s(t)$ . The rate of disease-induced deaths due to pulmonary tuberculosis for individuals in the symptomatic infectious class is given by  $\delta_1$ , whereas the rate of death due to the disease for symptomatic infectious individuals undergoing treatment is given by  $\delta_2$ . Treatment for different forms of pulmonary tuberculosis is assumed to be successful, and thus latent infected individuals undergoing treatment, asymptomatic infectious individuals undergoing treatment, and symptomatic infectious individuals undergoing treatment recover at rates of  $\xi_1, \xi_2$ , and  $\xi_3$ , respectively, and move to the recovered class R. The model assumes that recovered individuals become susceptible again after their immunity wanes at a rate of  $\sigma$ . The rate at which individuals die from causes other than pulmonary tuberculosis is denoted by  $\mu$ . It is worth noting that all parameters are positive constants.

The compartmental model illustrating the interaction of the human population in various classes is depicted in Figure 1.

The transmission model culminates in a ninedimensional system of ordinary differential equations as follows:

$$\begin{aligned} \frac{\mathrm{d}S}{\mathrm{d}t} &= (1-P)\pi + \sigma R - (\lambda + \mu)S, \\ \frac{\mathrm{d}V}{\mathrm{d}t} &= P\pi - ((1-\rho)\lambda + \mu)V, \\ \frac{\mathrm{d}E}{\mathrm{d}t} &= \lambda S + (1-\rho)\lambda V - (\chi_1 + \chi_2 + \theta_1 + \mu)E, \\ \frac{\mathrm{d}I_a}{\mathrm{d}t} &= \chi_1 E - (\theta_2 + \omega + \mu)I_a, \\ \frac{\mathrm{d}I_s}{\mathrm{d}t} &= \chi_2 E + \omega I_a - (\theta_3 + \delta_1 + \mu)I_s, \\ \frac{\mathrm{d}T_E}{\mathrm{d}t} &= \theta_1 E - (\xi_1 + \mu)T_E, \\ \frac{\mathrm{d}T_a}{\mathrm{d}t} &= \theta_2 I_a - (\xi_2 + \mu)T_a, \\ \frac{\mathrm{d}T_s}{\mathrm{d}t} &= \theta_3 I_s - (\delta_2 + \xi_3 + \mu)T_s, \\ \frac{\mathrm{d}R}{\mathrm{d}t} &= \xi_1 T_E + \xi_2 T_a + \xi_3 T_s - (\sigma + \mu)R. \end{aligned}$$
(1)

We assume that the model parameters are positive, and the initial conditions of system (1) are given as follows:  $S(0) \ge 0, V(0) \ge 0, E(0) \ge 0, I_a(0) \ge 0, I_s(0) \ge 0, T_E(0) \ge 0,$  $T_a(0) \ge 0, T_s(0) \ge 0,$  and  $R(0) \ge 0.$ 

#### 3. Model Analysis

*3.1. Invariant Region.* The model system (1) deals with the human population, and thus we need to demonstrate that its solutions are bounded for all time t > 0.

**Theorem 1.** *Given the positive initial conditions, the feasible region is defined as follows:* 

$$\Omega = \left\{ \left( S(t), V(t), E(t), I_a(t), I_s(t), T_E(t), T_a(t), T_s(t), R(t) \right) \in \mathbb{R}^9_+ : N \le \frac{\pi}{\mu} \right\}$$
(2)

*Proof.* The sum of all equations in system (1) represents the total human population in the model and satisfies the following equation:

$$\frac{\mathrm{d}N}{\mathrm{d}t} = \pi - \mu N - \delta_1 I_s - \delta_2 T_s \le \pi - \mu N$$

$$\Longrightarrow \frac{\mathrm{d}N}{\mathrm{d}t} \le \pi - \mu N.$$
(3)

By integrating inequality (3) and applying the initial conditions, we obtain

$$N(t) \le \frac{\pi}{\mu} + \left( N(0) - \frac{\pi}{\mu} \right) e^{-\mu t}.$$
 (4)

As  $t \longrightarrow \infty$  in inequality (3), the population  $N(t) \longrightarrow \pi/\mu$ , implying that  $0 \le N(t) \le \pi/\mu$ . Thus, the feasible solution of the system enters and remains in the region:



FIGURE 1: Schematic model diagram for the propagation and control of pulmonary tuberculosis disease in Kenya.

$$\Omega = \left\{ \left( S(t), V(t), E(t), I_a(t), I_s(t), T_E(t), T_a(t), T_s(t), R(t) \right) \in \mathbb{R}^9_+ : N \le \frac{\pi}{\mu} \right\}.$$
(5)

Therefore, the basic model is well-posed epidemiologically and mathematically, and hence it is sufficient to study its dynamics in  $\Omega$ .

3.2. The Basic Reproduction Number and the Control Reproduction Number. The basic reproduction number is a threshold parameter that governs the spread of disease. It is defined as the average number of secondary infections caused by a single infectious individual during his entire infectious period in a population that is entirely susceptible [25].

The control reproduction number is defined as the expected number of secondary infections produced by an index-infected individual in a population that is not entirely susceptible due to the presence of control measures [26].

In the absence of pulmonary tuberculosis,  $E = I_a = I_s = T_E = T_a = T_s = R = 0$ . Therefore, system (1) has a disease-free equilibrium given by

$$B^{0} = \left(S^{0}, V^{0}, E^{0}, I_{a}^{0}, I_{s}^{0}, T_{E}^{0}, T_{a}^{0}, T_{s}^{0}, R^{0}\right)$$
$$= \left(\frac{(1-P)\pi}{\mu}, \frac{P\pi}{\mu}, 0, 0, 0, 0, 0, 0, 0, 0\right).$$
(6)

We use the next-generation matrix to obtain the control reproduction number as given by [27].

Let  $X = (E, I_a, I_s, T_a, T_s)^T$ , then it follows from system (1) that

$$\frac{\mathrm{d}X}{\mathrm{d}t} = f - \nu,\tag{7}$$

where f and v are matrices representing the new infections and transition terms, respectively, given as follows:

$$f = \begin{bmatrix} \lambda S + (1 - \rho)\lambda V \\ 0 \\ 0 \\ 0 \end{bmatrix},$$

$$v = \begin{bmatrix} (\chi_1 + \chi_2 + \theta_1 + \mu)E \\ (\theta_2 + \omega + \mu)I_a - \chi_1E \\ (\theta_3 + \delta_1 + \mu)I_S - \chi_2E - \omega I_a \\ (\xi_2 + \mu)T_a - \theta_2I_a \\ (\xi_3 + \delta_2 + \mu)T_S - \theta_3I_S \end{bmatrix}.$$
(8)

The Jacobian matrices of the new infections and transition terms at the disease-free equilibrium are given, respectively, as follows:

The dominant eigenvalue corresponding to the spectral radius  $\rho(\text{FV}^{-1})$  of the matrix  $\text{FV}^{-1}$  is the control reproduction number ( $R_{\text{cVST}}$ ) with vaccination, screening,

and treatment of all forms of pulmonary tuberculosis as the intervention strategies.  $R_{cVST}$  is given as follows:

$$R_{\text{cVST}} = \frac{\beta \left[ \left( \eta_1 \chi_1 x_3 + \chi_1 \omega + \chi_2 x_2 \right) x_5 x_6 + \eta_3 \theta_2 \chi_1 x_3 x_6 + \eta_2 \theta_3 x_5 \left( \chi_1 \omega + \chi_2 x_2 \right) \right] \left[ (1 - P)\pi/\mu + (1 - \rho)P\pi/\mu \right]}{x_1 x_2 x_3 x_5 x_6}, \tag{10}$$

$$\begin{aligned} x_1 &= (\chi_1 + \chi_2 + \theta_1 + \mu), \\ x_2 &= (\theta_2 + \omega + \mu), \\ x_3 &= (\theta_3 + \delta_1 + \mu), \\ x_5 &= (\xi_2 + \mu), \\ x_6 &= (\delta_2 + \xi_3 + \mu). \end{aligned}$$
 (11)

Without vaccination intervention, the fraction of recruits vaccinated, *P*, equals zero, and consequently, the parameter,  $\rho$ , for vaccine efficacy becomes zero since there will be no vaccinated population. Substituting  $P = \rho = 0$  in (10) gives the control reproduction number ( $R_{cST}$ ) with screening and treatment as the only intervention strategies.  $R_{cST}$  is thus given as follows:

$$R_{\rm cST} = \frac{\beta [(\eta_1 \chi_1 x_3 + \chi_1 \omega + \chi_2 x_2) x_5 x_6 + \eta_3 \theta_2 \chi_1 x_3 x_6 + \eta_2 \theta_3 x_5 (\chi_1 \omega + \chi_2 x_2)] \pi/\mu}{x_1 x_2 x_3 x_5 x_6}.$$
(12)

Considering the presence of latent infected and asymptomatic infectious individuals in the population without their screening, the parameters  $\theta_1$ ,  $\theta_2$ , and  $\xi_2$  become zero. Substituting  $\theta_1 = \theta_2 = \xi_2 = 0$  in (10) gives the

control reproduction number  $(R_{\rm cVT_s})$  with vaccination and treatment of the symptomatic infectious population as the only intervention strategies.  $R_{\rm cVT_s}$  is thus given as follows:

$$R_{\rm cVT_s} = \frac{\beta \left[ \left[ \eta_1 \chi_1 \left( \theta_3 + \delta_1 + \mu \right) + \chi_1 \omega + \chi_2 \left( \mu + \omega \right) \right] \left( \xi_3 + \delta_2 + \mu \right) + \eta_2 \theta_3 \left[ \chi_1 \omega + \chi_2 \left( \mu + \omega \right) \right] \right] \left[ (1 - P)\pi/\mu + (1 - \rho)P\pi/\mu \right]}{\left( \chi_1 + \chi_2 + \mu \right) \left( \omega + \mu \right) \left( \theta_3 + \delta_1 + \mu \right) \left( \delta_2 + \xi_3 + \mu \right)}.$$
 (13)

When there is no vaccination of recruits and screening of both latent infected and asymptomatic infectious populations, the parameters  $\theta_1$ ,  $\theta_2$ ,  $\xi_2$ , *P*, and  $\rho$  become zero. Substituting  $\theta_1 = \theta_2 = \xi_2 = P = \rho = 0$  in (10) gives the control reproduction number  $(R_{cT_s})$  with the treatment of the symptomatic infectious population as the only intervention strategy.  $R_{cT_s}$  is thus given as follows:

$$R_{cT_{s}} = \frac{\beta \left[ \left[ \eta_{1} \chi_{1} \left( \theta_{3} + \delta_{1} + \mu \right) + \chi_{1} \omega + \chi_{2} \left( \mu + \omega \right) \right] \left( \xi_{3} + \delta_{2} + \mu \right) + \eta_{2} \theta_{3} \left[ \chi_{1} \omega + \chi_{2} \left( \mu + \omega \right) \right] \right] \pi / \mu}{\left( \chi_{1} + \chi_{2} + \mu \right) \left( \omega + \mu \right) \left( \theta_{3} + \delta_{1} + \mu \right) \left( \delta_{2} + \xi_{3} + \mu \right)}.$$
(14)

Considering no intervention measures in place, that is, when there is no vaccination of the recruits, screening of both latent and asymptomatic infectious populations, and treatment of all forms of pulmonary tuberculosis, the parameters P,  $\rho$ ,  $\theta_1$ ,  $\theta_2$ ,  $\theta_3$ ,  $\xi_2$ ,  $\xi_3$ , and  $\delta_2$  become zero. Substituting  $P = \rho = \theta_1 = \theta_2 = \theta_3 = \xi_2 = \xi_3 = \delta_2 = 0$  in (10) gives the basic reproduction number ( $R_{\rho}$ ) given as follows:

$$R_{o} = \frac{\beta [\eta_{1} \chi_{1} (\delta_{1} + \mu) + \chi_{1} \omega + \chi_{2} (\mu + \omega)] \pi / \mu}{(\chi_{1} + \chi_{3} + \mu) (\omega + \mu) (\delta_{1} + \mu)}.$$
 (15)

#### 3.3. Local Stability of Disease-Free Equilibrium

**Theorem 2.** The disease-free equilibrium point is locally asymptotically stable if  $R_{\text{cVST}} < 1$  and unstable if  $R_{\text{cVST}} > 1$ .

*Proof.* To prove the local stability of the disease-free equilibrium, we obtain the Jacobian matrix of the system at the disease-free equilibrium as follows:

$$J(B^{0}) = \begin{bmatrix} -\mu & 0 & 0 & -\beta\eta_{1}S^{0} & -\beta S^{0} & 0 & -\beta\eta_{3}S^{0} & -\beta\eta_{2}S^{0} & \sigma \\ 0 & -\mu & 0 & -H_{1} & -H_{2} & 0 & -H_{3} & -H_{4} & 0 \\ 0 & 0 & -x_{1} & H_{5} & H_{6} & 0 & H_{7} & H_{8} & 0 \\ 0 & 0 & \chi_{1} & -x_{2} & 0 & 0 & 0 & 0 \\ 0 & 0 & \chi_{2} & \omega & -x_{3} & 0 & 0 & 0 \\ 0 & 0 & \theta_{1} & 0 & 0 & -x_{4} & 0 & 0 \\ 0 & 0 & \theta_{1} & 0 & 0 & -x_{5} & 0 & 0 \\ 0 & 0 & 0 & \theta_{2} & 0 & 0 & -x_{5} & 0 & 0 \\ 0 & 0 & 0 & 0 & \theta_{3} & 0 & 0 & -x_{6} & 0 \\ 0 & 0 & 0 & 0 & 0 & \xi_{1} & \xi_{2} & \xi_{3} & -x_{7} \end{bmatrix},$$
(16)

where

$$\begin{aligned} x_{1} &= (\chi_{1} + \chi_{2} + \theta_{1} + \mu), \\ x_{2} &= (\theta_{2} + \omega + \mu), \\ x_{3} &= (\theta_{3} + \delta_{1} + \mu), \\ x_{4} &= (\xi_{1} + \mu), \\ x_{5} &= (\xi_{2} + \mu), \\ x_{5} &= (\xi_{2} + \mu), \\ x_{7} &= (\sigma + \mu), \\ H_{1} &= (1 - \rho)\beta\eta_{1}V^{0}, \\ H_{2} &= (1 - \rho)\beta V^{0}, \\ H_{3} &= (1 - \rho)\beta\eta_{2}V^{0}, \\ H_{4} &= (1 - \rho)\beta\eta_{2}V^{0}, \\ H_{5} &= \beta\eta_{1} \left[ S^{0} + (1 - \rho)V^{0} \right], \\ H_{6} &= \beta \left[ S^{0} + (1 - \rho)V^{0} \right], \\ H_{7} &= \beta\eta_{3} \left[ S^{0} + (1 - \rho)V^{0} \right], \\ H_{8} &= \beta\eta_{2} \left[ S^{0} + (1 - \rho)V^{0} \right]. \end{aligned}$$
(17)

Evaluating the eigenvalues of (16), we obtain

$$\{-(\mu+\lambda)\}\{-(\mu+\lambda)\}\{-(x_{7}+\lambda)\} \begin{cases} -x_{1}-\lambda & H_{5} & H_{6} & 0 & H_{7} & H_{8} \\ \chi_{1} & -x_{2}-\lambda & 0 & 0 & 0 \\ \chi_{2} & \omega & -x_{3}-\lambda & 0 & 0 & 0 \\ \theta_{1} & 0 & 0 & -x_{4}-\lambda & 0 & 0 \\ 0 & \theta_{2} & 0 & 0 & -x_{5}-\lambda & 0 \\ 0 & 0 & \theta_{3} & 0 & 0 & -x_{6}-\lambda \end{cases} = 0.$$

$$(18)$$

or

Either

$$\{-(\mu+\lambda)\}\{-(\mu+\lambda)\}\{-(x_7+\lambda)\}=0,$$
 (19)

$$\begin{vmatrix} -x_1 - \lambda & H_5 & H_6 & 0 & H_7 & H_8 \\ \chi_1 & -x_2 - \lambda & 0 & 0 & 0 & 0 \\ \chi_2 & \omega & -x_3 - \lambda & 0 & 0 & 0 \\ \theta_1 & 0 & 0 & -x_4 - \lambda & 0 & 0 \\ 0 & \theta_2 & 0 & 0 & -x_5 - \lambda & 0 \\ 0 & 0 & \theta_3 & 0 & 0 & -x_6 - \lambda \end{vmatrix} = 0.$$

$$(20)$$

Simplifying (19), we have

$$(\mu + \lambda)(\mu + \lambda)(x_7 + \lambda) = 0.$$
(21)

By the Routh-Hurwitz criteria, (21) has strictly negative roots given as follows:

$$\lambda_1 = \lambda_2$$
  
=  $-\mu < 0,$  (22)  
 $\lambda_3 = -x_7 < 0.$ 

The characteristic polynomial of (20) is obtained as follows:

$$M_1 \lambda^6 + M_2 \lambda^5 + M_3 \lambda^4 + M_4 \lambda^3 + M_5 \lambda^2 + M_6 \lambda + M_7 = 0,$$
(23)

where  $M_1, M_2, M_3, M_4, M_5, M_6$ , and  $M_7$  are determined using Mathematica software as follows:

$$\begin{split} M_{1} &= 1 > 0, \\ M_{2} &= x_{1} + x_{2} + x_{3} + x_{4} + x_{5} + x_{6}, \\ M_{3} &= x_{6} \left(x_{1} + x_{2} + x_{3} + x_{4} + x_{5}\right) + x_{5} \left(x_{1} + x_{2} + x_{4}\right) + x_{4} \left(x_{1} + x_{2} + x_{3}\right) + x_{3} \left(x_{1} + x_{2}\right) + x_{1} x_{2} + H_{5} \chi_{1} - H_{6} \chi_{2}, \\ M_{4} &= x_{6} \left[x_{5} \left(x_{1} + x_{2} + x_{3} + x_{4}\right) + x_{4} \left(x_{1} + x_{2} + x_{3}\right) + x_{3} \left(x_{1} + x_{2}\right) + x_{1} x_{2} - (H_{5} \chi_{1} + H_{6} \chi_{2})\right] \\ &+ x_{5} \left[x_{4} \left(x_{1} + x_{2} + x_{3}\right) + x_{3} \left(x_{1} + x_{2}\right) + x_{1} x_{2} - (H_{5} \chi_{1} + H_{6} \chi_{2})\right] + x_{4} \left[x_{3} \left(x_{1} + x_{2}\right) + x_{1} x_{2} - (H_{5} \chi_{1} + H_{5} \chi_{2})\right] \\ &+ x_{3} \left(x_{1} x_{2} - H_{5} \chi_{1}\right) - x_{2} H_{6} \chi_{2} - \omega H_{6} \chi_{1} - \theta_{2} H_{7} \chi_{1} - \theta_{3} H_{8} \chi_{2}, \\ M_{5} &= x_{6} \left[x_{4} x_{5} \left(x_{1} + x_{2} + x_{3}\right) + x_{3} x_{5} \left(x_{1} + x_{2}\right) + x_{2} x_{5} x_{1} + x_{3} x_{4} \left(x_{1} + x_{2}\right) + x_{1} x_{2} \left(x_{3} + x_{4}\right) - H_{5} \chi_{1} \left(x_{3} + x_{4} + x_{5}\right) \right) \\ &- H_{6} \omega \chi_{1} - H_{6} \chi_{2} \left(x_{2} + x_{4} + x_{5}\right) - H_{7} \theta_{2} \chi_{1}\right] + x_{5} \left[x_{1} x_{2} \left(x_{3} + x_{4}\right) + x_{3} x_{4} \left(x_{1} + x_{2}\right) - H_{5} \chi_{1} \left(x_{3} + x_{4}\right) - H_{6} \omega \chi_{1} \right) \\ &- H_{6} \chi_{2} \left(x_{2} + x_{4}\right) - H_{8} \theta_{3} \chi_{2}\right] + x_{4} \left[x_{3} x_{2} x_{1} - H_{5} \chi_{1} x_{3} - H_{6} \omega \chi_{1} - H_{6} \chi_{2} x_{2} - H_{7} \theta_{2} \chi_{1} - H_{8} \theta_{3} \chi_{2}\right] \\ &- H_{7} \theta_{2} \chi_{1} x_{3} - H_{8} \theta_{3} \chi_{2} x_{2} - H_{8} \theta_{3} \omega \chi_{1}, \\ M_{6} &= x_{6} \left[x_{1} x_{2} \left(x_{3} x_{4} + x_{3} x_{5} + x_{4} x_{5}\right) - H_{7} \theta_{2} \chi_{1} \left(x_{3} + x_{4}\right)\right] + x_{5} \left[x_{1} x_{2} x_{3} x_{4} - H_{5} \chi_{1} x_{3} x_{4} - H_{6} \chi_{2} x_{4} x_{5} \right) \\ &- H_{6} \omega \chi_{1} x_{4} - H_{8} \theta_{3} \chi_{2} \left(x_{2} + x_{4}\right) - H_{8} \omega \theta_{3} \chi_{1}\right] + x_{4} \left[-H_{7} \theta_{2} \chi_{1} x_{3} - H_{8} \theta_{3} \left(x_{2} \chi_{2} + \chi_{1} \omega\right)\right]. \\ M_{7} &= x_{1} x_{2} x_{3} x_{4} x_{5} x_{6} - x_{4} \left[x_{5} x_{6} \left[x_{3} H_{5} \chi_{1} + H_{6} \left(x_{2} x_{2} + \chi_{1} \omega\right)\right] + H_{7} \chi_{1} \theta_{2} x_{3} x_{6} + H_{8} \theta_{3} x_{5} \left(\chi_{2} x_{2} + \chi_{1} \omega\right)\right]. \\ \end{pmatrix}$$

By the Routh-Hurwitz criteria,

From  $M_7 > 0$ , we have

$$M_1 > 0, M_3 > 0, M_5 > 0 \text{ and } M_7 > 0.$$
 (25)

$$x_{1}x_{2}x_{3}x_{4}x_{5}x_{6} - x_{4}[x_{5}x_{6}[x_{3}H_{5}\chi_{1} + H_{6}(\chi_{2}x_{2} + \chi_{1}\omega)] + H_{7}\chi_{1}\theta_{2}x_{3}x_{6} + H_{8}\theta_{3}x_{5}(\chi_{2}x_{2} + \chi_{1}\omega)] > 0$$

$$\implies [x_{5}x_{6}[x_{3}H_{5}\chi_{1} + H_{6}(\chi_{2}x_{2} + \chi_{1}\omega)] + H_{7}\chi_{1}\theta_{2}x_{3}x_{6} + H_{8}\theta_{3}x_{5}(\chi_{2}x_{2} + \chi_{1}\omega)] < x_{1}x_{2}x_{3}x_{5}x_{6}$$

$$\implies \frac{[x_{5}x_{6}[x_{3}H_{5}\chi_{1} + H_{6}(\chi_{2}x_{2} + \chi_{1}\omega)] + H_{7}\chi_{1}\theta_{2}x_{3}x_{6} + H_{8}\theta_{3}x_{5}(\chi_{2}x_{2} + \chi_{1}\omega)] < x_{1}x_{2}x_{3}x_{5}x_{6}}{x_{1}x_{2}x_{3}x_{5}x_{6}}$$

$$\implies \frac{[x_{5}x_{6}[x_{3}H_{5}\chi_{1} + H_{6}(\chi_{2}x_{2} + \chi_{1}\omega)] + H_{7}\chi_{1}\theta_{2}x_{3}x_{6} + H_{8}\theta_{3}x_{5}(\chi_{2}x_{2} + \chi_{1}\omega)] < x_{1}x_{2}x_{3}x_{5}x_{6}}{x_{1}x_{2}x_{3}x_{5}x_{6}}$$

$$\implies (26)$$

Substituting  $H_5 = \beta \eta_1 [S^0 + (1 - \rho)V^0]$ ,  $H_6 = \beta [S^0 + (1 - \rho)V^0]$ ,  $H_7 = \beta \eta_3 [S^0 + (1 - \rho)V^0]$ ,  $H_8 = \beta \eta_2 [S^0 + (1 - \rho)V^0]$ ,  $S^0 = (1 - P)\pi/\mu$ , and  $V^0 = P\pi/\mu$  in inequality (26) and rearranging yield

$$\frac{\beta[(\eta_1\chi_1x_3 + \chi_1\omega + \chi_2x_2)x_5x_6 + \eta_3\theta_2\chi_1x_3x_6 + \eta_2\theta_3x_5(\chi_1\omega + \chi_2x_2)][(1-P)\pi/\mu + (1-\rho)P\pi/\mu]}{x_1x_2x_3x_5x_6} < 1.$$
(27)

However,

$$\frac{\beta[(\eta_1\chi_1x_3 + \chi_1\omega + \chi_2x_2)x_5x_6 + \eta_3\theta_2\chi_1x_3x_6 + \eta_2\theta_3x_5(\chi_1\omega + \chi_2x_2)][(1-P)\pi/\mu + (1-\rho)P\pi/\mu]}{x_1x_2x_3x_5x_6} = R_{\text{cVST}}.$$
(28)

Therefore, (27) can be expressed as follows:

$$R_{\rm cVST} < 1.$$
 (29)

From (29), the disease-free equilibrium is locally asymptotically stable if  $R_{\text{cVST}} < 1$ . This implies that each infectious individual infects, on average, less than one susceptible person during the infectious period, resulting in the disease dying out [27].

3.4. The Endemic Equilibrium. The endemic equilibrium  $B^* = (S^*, V^*, E^*, I_a^*, I_s^*, T_E^*, T_a^*, T_s^*R^*)$  is evaluated by equating the model system of (1) to zero. The steady-state solution for the model equations is as follows:

$$S^{*} = \frac{(1-P)\pi + \sigma R^{*}}{\lambda^{**} + \mu},$$

$$V^{*} = \frac{P\pi}{(1-\rho)\lambda^{**} + \mu'},$$

$$E^{*} = \frac{\lambda^{**}(1-P)\pi + \lambda^{**}\sigma R^{*} + (1-\rho)\lambda^{**}P\pi}{(\lambda^{**} + \mu)(\chi_{1} + \chi_{2} + \theta_{1} + \mu)((1-\rho)\lambda^{**} + \mu)},$$

$$I_{a}^{*} = \frac{\chi_{1}[\lambda^{**}(1-P)\pi + \lambda^{**}\sigma R^{*} + (1-\rho)\lambda^{**}P\pi]}{(\lambda^{**} + \mu)(\chi_{1} + \chi_{2} + \theta_{1} + \mu)((1-\rho)\lambda^{**} + \mu)(\theta_{2} + \omega + \mu)},$$

$$I_{s}^{*} = \frac{[\chi_{2}(\theta_{2} + \omega + \mu) + \omega\chi_{1}][\lambda^{**}(1-P)\pi + \lambda^{**}\sigma R^{*} + (1-\rho)\lambda^{**}P\pi]}{(\theta_{3} + \delta_{1} + \mu)(\lambda^{**} + \mu)(\chi_{1} + \chi_{2} + \theta_{1} + \mu)((1-\rho)\lambda^{**} + \mu)(\theta_{2} + \omega + \mu)},$$

$$T_{E}^{*} = \frac{\theta_{1}[\lambda^{**}(1-P)\pi + \lambda^{**}\sigma R^{*} + (1-\rho)\lambda^{**}P\pi]}{(\xi_{1} + \mu)(\lambda^{**} + \mu)(\chi_{1} + \chi_{2} + \theta_{1} + \mu)((1-\rho)\lambda^{**} + \mu)},$$

$$T_{a}^{*} = \frac{\theta_{2}\chi_{1}[\lambda^{**}(1-P)\pi + \lambda^{**}\sigma R^{*} + (1-\rho)\lambda^{**}P\pi]}{(\lambda^{**} + \mu)(\chi_{1} + \chi_{2} + \theta_{1} + \mu)((1-\rho)\lambda^{**} + \mu)(\xi_{2} + \omega + \mu)(\xi_{2} + \mu)},$$

$$T_{s}^{*} = \frac{[\chi_{2}(\theta_{2} + \omega + \mu) + \omega\chi_{1}]\theta_{3}[\lambda^{**}(1-P)\pi + \lambda^{**}\sigma R^{*} + (1-\rho)\lambda^{**}P\pi]}{(\theta_{3} + \delta_{1} + \mu)(\lambda^{**} + \mu)(\chi_{1} + \chi_{2} + \theta_{1} + \mu)((1-\rho)\lambda^{**} + \mu)(\theta_{2} + \omega + \mu)(\theta_{2} + \xi_{3} + \mu)},$$

$$R^{*} = \frac{[\lambda^{**}S^{*} + (1-\rho)\lambda^{**}P\pi][D_{1} + D_{2} + D_{3}]}{(\lambda^{**} + \mu)((1-\rho)\lambda^{**} + \mu)(\chi_{1} + \chi_{2} + \theta_{1} + \mu)(\theta_{2} + \omega + \mu)(\theta_{3} + \delta_{1} + \mu)A_{1}A_{2}A_{3}A_{4}},$$

where

$$\begin{split} D_1 &= \xi_1 \theta_1 \left( \theta_3 + \delta_1 + \mu \right) \left( \theta_2 + \omega + \mu \right) \left( \delta_2 + \xi_3 + \mu \right) \left( \xi_2 + \mu \right), \\ D_2 &= \xi_2 \theta_2 \chi_1 \left( \theta_3 + \delta_1 + \mu \right) \left( \delta_2 + \xi_3 + \mu \right) \left( \xi_1 + \mu \right), \\ D_3 &= \xi_3 \theta_3 \left[ \chi_2 \left( \theta_2 + \omega + \mu \right) + \omega \chi_1 \right] \left( \xi_1 + \mu \right) \left( \xi_2 + \mu \right), \\ A_1 &= \left( \xi_1 + \mu \right), \\ A_2 &= \left( \xi_2 + \mu \right), \\ A_3 &= \left( \delta_2 + \xi_3 + \mu \right), \\ A_4 &= (\sigma + \mu). \end{split}$$

3.4.1. Global Stability of Endemic Equilibrium Point. We consider the method of the Lyapunov function to prove the global stability of the endemic equilibrium point. We propose a logarithmic Lyapunov function L defined by

$$L(S^{*}, V^{*}, E^{*}, I_{a}^{*}, I_{s}^{*}, T_{E}^{*}, T_{a}^{*}, T_{s}^{*}, R^{*}) = \left(S - S^{*} + S^{*} \ln \frac{S^{*}}{S}\right) + \left(V - V^{*} + V^{*} \ln \frac{V^{*}}{V}\right) + \left(E - E^{*} + E^{*} \ln \frac{E^{*}}{E}\right) \\ + \left(I_{a} - I_{a}^{*} + I_{a}^{*} \ln \frac{I_{a}^{*}}{I_{a}}\right) + \left(I_{s} - I_{s}^{*} + I_{s}^{*} \ln \frac{I_{s}^{*}}{I_{s}}\right) + \left(T_{E} - T_{E}^{*} + T_{E}^{*} \ln \frac{T_{E}^{*}}{T_{E}}\right) \\ + \left(T_{a} - T_{a}^{*} + T_{a}^{*} \ln \frac{T_{a}^{*}}{T_{a}}\right) + \left(T_{s} - T_{s}^{*} + T_{s}^{*} \ln \frac{T_{s}^{*}}{T_{s}}\right) + \left(R - R^{*} + R^{*} \ln \frac{R^{*}}{R}\right).$$
(32)

(31)

The derivatives of L along the solution of the model system (1) give

$$\frac{dL}{dt} = \left(\frac{S-S^*}{S}\right) \frac{dS}{dt} + \left(\frac{V-V^*}{V}\right) \frac{dV}{dt} + \left(\frac{E-E^*}{E}\right) \frac{dE}{dt} + \left(\frac{I_a - I_a^*}{I_a}\right) \frac{dI_a}{dt} + \left(\frac{I_s - I_s^*}{I_s}\right) \frac{dI_s}{dt} + \left(\frac{T_e - T_E^*}{T_E}\right) \frac{dT_E}{dt} + \left(\frac{T_a - T_a^*}{T_a}\right) \frac{dT_a}{dt} + \left(\frac{T_s - T_s^*}{T_s}\right) \frac{dT_s}{dt} + \left(\frac{R-R^*}{R}\right) \frac{dR}{dt}.$$
(33)  
e expressions dS/dt, dV/dt, dE/dt, dI\_a/dt,  $\frac{dL}{dt} = Q - W$ , (34)

Substituting the expressions dS/dt, dV/dt, dE/dt, d $I_a$ /dt, d $I_s$ /dt, d $T_E$ /dt, d $T_a$ /dt, d $T_s$ /dt, and dR/dt from the model system (1) into (33) and simplifying give

where

$$\begin{split} Q &= \pi + \sigma R + \frac{\sigma R^* S^*}{S} + \frac{\lambda S^* E^*}{E} (1 - \rho)\lambda V + \frac{(1 - \rho)\lambda V^* E^*}{E} + \chi_1 E + \frac{\chi_1 I_a^* E^*}{I_a} + \chi_2 E \\ &+ \frac{\chi_2 I_s^* E^*}{I_s} + \omega I_a + \frac{\omega I_s^* I_a^*}{I_s} + \theta_1 E + \frac{\theta_1 T_E^* E^*}{T_E} + \theta_2 I_a + \frac{\theta_2 I_a^* T_a^*}{T_a} + \theta_3 T_s + \frac{\theta_3 I_s^* T_s^*}{T_s} \\ &+ \xi_1 T_E + \frac{\xi_1 T_E^* R^*}{R} + \xi_2 T_a + \frac{\xi_2 T_a^* R^*}{R} + \xi_3 T_s + \frac{\xi_3 T_s^* R^*}{R} + \frac{\pi P V^*}{V} + \lambda S, \end{split}$$
$$W &= \frac{\pi S^*}{S} + \frac{\pi P S^*}{S} + \frac{\sigma R S^*}{S} + \frac{\lambda S E^*}{E} + \frac{\pi P V^*}{V} + (1 - \rho)\lambda V^* + \frac{(1 - \rho)\lambda E^*}{E^*} + \chi_1 E^* + \frac{\chi_1 E I_a^*}{I_a} + \chi_2 E^* \end{split}$$

$$+ \chi_{2}EI_{s}^{*} + \omega I_{a}^{*} + \frac{\omega I_{a}I_{s}^{*}}{I_{s}} + \theta_{1}E^{*} + \frac{\theta_{1}ET_{E}^{*}}{T_{E}} + \theta_{2}I_{a}^{*} + \frac{\theta_{2}I_{a}T_{a}^{*}}{T_{a}} + \theta_{3}I_{s}^{*} + \frac{\theta_{3}I_{s}T_{s}^{*}}{T_{s}} + \xi_{1}T_{E}^{*}$$

$$+ \frac{\xi_{1}T_{E}R^{*}}{R} + \xi_{2}T_{a}^{*} + \frac{\xi_{2}T_{a}R^{*}}{R} + \xi_{3}T_{s}^{*} + \frac{\xi_{2}T_{s}R^{*}}{R} + \lambda S^{*}$$

$$+ \frac{(S-S^{*})^{2}}{S}[\lambda + \mu] + \frac{(V-V^{*})^{2}}{V}[(1-\rho)\lambda + \mu] + \frac{(E-E^{*})^{2}}{E}[\chi_{1} + \chi_{2} + \theta_{1} + \mu]$$

$$+ \frac{(I_{a}-I_{a}^{*})^{2}}{I_{a}}[\theta_{2} + \omega + \mu] + \frac{(I_{s}-I_{s}^{*})^{2}}{I_{s}}[\theta_{3} + \delta_{1} + \mu]$$

$$+ \frac{(T_{E}-T_{E}^{*})^{2}}{T_{E}}[\xi_{1} + \mu] + \frac{(T_{a}-T_{a}^{*})}{T_{a}}[\xi_{2} + \mu] + \frac{(T_{s}-T_{s}^{*})^{2}}{T_{s}}[\delta_{2} + \xi_{3} + \mu] + \frac{(R-R^{*})^{2}}{R}[\sigma + \mu].$$

$$(35)$$

If  $Q \le W$ , then  $dL/dt \le 0$  and dL/dt = 0 if and only if  $S = S^*, V = V^*, E = E^*, I_a = I_a^*, I_s = I_s^*, T_E = T_E^*, T_a = T_a^*, T_s = T_s^*, R = R^*$ .

Therefore, the largest compact invariant set in  $\{(S^*, V^*, E^*, I_a^*, I_s^*, T_a^*, T_s^*, R^*) \in \Omega : dL/dt = 0\}$  is the singleton endemic equilibrium point  $B^*$ . Thus, from LaSalle's invariance principle [28], we conclude that as  $t \longrightarrow \infty$ , the solution of the model system (1) approaches the endemic equilibrium  $B^*$  when the control reproduction number  $R_{cVST}^* > 1$ . Therefore, the endemic equilibrium point  $B^*$  is globally asymptotically stable in the invariant set  $\Omega$  if Q < W.

3.5. Sensitivity Analysis on Control Reproduction Numbers. In this section, we present a sensitivity analysis of the reproduction numbers to determine the relative importance of the various parameters responsible for pulmonary tuberculosis transmission and prevalence in the population. We employed the normalized forward sensitivity index for this model as used by [29]. The normalized sensitivity index which measures the relative change in a parameter K with respect to the reproduction number  $R_c$  is given by

$$\Lambda_K^{R_c} = \frac{\partial R_c}{\partial K} \times \frac{K}{R_c}.$$
(36)

The parameter values in Table 1 are used to calculate the sensitivity indices of the reproduction numbers for the parameters  $\beta$ , *P*,  $\chi_1$ ,  $\chi_2$ ,  $\theta_1$ ,  $\theta_2$ , and  $\theta_3$ .

The calculated sensitivity indices of reproduction numbers are given in Table 2. A positive sensitivity index shows that the reproduction number is an increasing function of the corresponding parameter whereas a negative sensitivity index shows that the reproduction number is a decreasing function of the corresponding parameter. Thus, increasing a parameter with a positive sensitivity index holding other parameters constant increases the reproduction number whereas increasing a parameter with a negative sensitivity index while other factors are held constant decreases the reproduction number [37].

From Table 2, it is observed that  $\beta$ ,  $\chi_1$ , and  $\chi_2$  have positive sensitivity index values, and thus an increase in these parameters results in a corresponding increase in the

number of the infected population. On the other hand,  $\theta_1$ ,  $\theta_2$ ,  $\theta_3$ , and *P* have negative sensitivity index values, and thus an increase in these parameters results in a decrease in the number of infected populations. For instance, if the transmission rate,  $\beta$ , is increased by 10%, the reproduction numbers increase by 10%. On the other hand, increasing screening rate of latent infected,  $\theta_1$ , by 10% decreases  $R_{cVST}$  by 6.673668% while increasing screening rate of asymptomatic infectious,  $\theta_2$ , by 10% decreases  $R_{cVST}$  by 0.751639%.

#### 4. Numerical Simulations of the Model

Numerical simulations of the model system (1) have been done to explore pulmonary tuberculosis epidemic behavior. The simulations were carried out in MATLAB ordinary differential equations inbuilt solver, ode45 function. Numerical simulations have been performed using data from the Kenyan population. The baseline parameters used for numerical simulations are given in Table 1. The initial values states are gives as S(0) = 1182969, V(0) = 30920683, E(0) = 17923767,  $I_a(0) = 127595$ ,  $I_s(0) = 200000$ ,  $T_E(0) = 500000$ ,  $T_a(0) = 50000$ , and  $T_s(0) = 113155$ , R(0) = 2753131 as obtained from published Kenyan data [30–34]. The simulation results are presented graphically as shown in Figures 2–7.

4.1. Effects of Various Intervention Strategies on the Control Reproduction Number. Equations (10), (12)-(14) represent the control reproduction numbers with various intervention strategies:  $R_{\rm cVST}$  for vaccination, screening, and treatment of all infected cases; R<sub>cST</sub> for screening and treatment of all infected cases;  $R_{\rm cVT_s}$  for vaccination and treatment of symptomatic infectious population; and  $R_{cT_s}$  for treatment of symptomatic infectious population alone. Figure 2 illustrates the effects of varying reproduction numbers with respect to transmission rate,  $\beta$ . It is observed that  $R_{\rm cVST} < R_{\rm cST} < R_{\rm cVT_S} < R_{\rm cT_S}$ . This indicates that a combination of vaccination, screening, and treatment of all infected cases is the most effective control measure in reducing infection transmission in the population. Following this, a combination of screening and treatment of all infected cases proves effective, while the combination of vaccination

Parameter	Description	Value	Reference
$\eta_3$	Transmission coefficient for asymptomatic individuals undergoing treatment	0.00000126	Estimated
$\eta_2$	Transmission coefficient for symptomatic individuals undergoing treatment	0.00002	Estimated
$\eta_1$	Transmission coefficient for the asymptomatic infectious individuals	0.0003	Estimated
σ	The rate at which the immunity of recovered wanes	$0.003 \text{ year}^{-1}$	[30]
μ	Natural death rate	0.0147 year <sup>-1</sup>	[31]
π	Recruitment rate	0.021 year <sup>-1</sup>	[32]
$\chi_1$	Progression rate of latent infected to asymptomatic individuals	0.05 year <sup>-1</sup>	[33]
$\delta_2$	Rate of tuberculosis disease-induced deaths during treatment	$0.065 \text{ year}^{-1}$	[30]
$\chi_2$	Progression rate of latent infected to infectious individuals with disease symptoms	$0.1 \text{ year}^{-1}$	[33]
β	Transmission rate of pulmonary tuberculosis	$0.15 \text{ year}^{-1}$	[30]
$\theta_1$	Rate of screening latently infected individuals	$0.34 \text{ year}^{-1}$	[34]
ρ	Vaccine efficacy	0.5	[11]
$\delta_1$	Rate of tuberculosis disease-induced deaths	$0.5 \text{ year}^{-1}$	[34]
$\theta_2$	Rate of screening asymptomatic infectious individuals	$0.5 \text{ year}^{-1}$	[35]
ω	The rate at which asymptomatic infectious individuals exhibit symptoms of the disease	$0.6 \text{ year}^{-1}$	[35]
$\theta_3$	The rate at which symptomatic individuals are treated	$0.68 \text{ year}^{-1}$	[30]
ξ	The recovery rate of treated symptomatic individuals	$0.75 \text{ year}^{-1}$	[32]
P	Vaccination rate	$0.8 \text{ year}^{-1}$	[31]
$\xi_2$	The recovery rate of treated asymptomatic infectious individuals	$0.8 \text{ year}^{-1}$	[30]
$\bar{\xi_1}$	The recovery rate of latent treated	0.85 year <sup>-1</sup>	[36]

TABLE 1: Baseline parameters used in simulation.

TABLE 2: Sensitivity indices of reproduction numbers with respect to some model parameters.

Parameter	Sensitivity index			
Sensitivity indices of $R_{\rm cVST}$				
$\theta_2$	-0.751639			
$ heta_1$	-0.673668			
Р	-0.666667			
$\theta_3$	-0.5690921			
$\chi_1$	+0.11309			
$\chi_2$	+0.589703			
β	+1			
Sensitivity indices of R <sub>cST</sub>				
$\theta_2$	-0.751362			
$\theta_1$	-0.673668			
$ heta_3$	-0.0519066			
$\chi_1$	+0.113028			
$\chi_2$	+0.589756			
β	+1			
Sensitivity indices of R <sub>cVT</sub>				
Р	-0.6666667			
$\theta_3$	-0.0519052			
$\chi_1$	+0.024444			
$\chi_2$	+0.0648092			
β	+1			
Sensitivity indices of R <sub>cT</sub>				
$ heta_3$	-0.569053			
$\chi_1$	+0.0245257			
$\chi_2$	+0.0647274			
β	+1			

and treatment of symptomatic infectious ranks as the third appropriate strategy. Furthermore, it is noted that treating symptomatic cases alone is the least effective strategy in reducing infection transmission in the population.

4.2. Effects of Varying Screening and Treatment Rates on the Control Reproduction Number. In the model flowchart depicted in Figure 1,  $\theta_1$  represents the rate of screening latent infected individuals, with Figure 3 illustrating the effects of screening the latent infected population on the control reproduction number. Similarly, according to the model flowchart shown in Figure 1,  $\theta_2$  represents the rate of screening the asymptomatic infectious population, with Figure 4 displaying the effects of screening this population on the control reproduction number. It is observed that increasing screening rates for both latent infected and asymptomatic infectious populations reduce the control reproduction number, consequently decreasing the rate of infection transmission. Screening and treating latent infections reduce reactivation, subsequently decreasing infection transmission in the population. The asymptomatic infectious population, experiencing no symptoms, continues their daily routines without seeking medical intervention. This behavior leads to more interactions with susceptible individuals, contributing to a high rate of infection transmission in the population. Therefore, it is prudent to screen and treat asymptomatic infectious individuals since they spread infections incessantly without being noticed.

In the model flowchart depicted in Figure 1,  $\theta_3$  represents the rate of treating the symptomatic infectious population, while Figure 5 shows the effects of varying treatment on the control reproduction number for this population. It is observed that increasing the treatment rate of the symptomatic infectious population reduces the control reproduction number. This reduction is attributed to the decrease in the population that is infectious, consequently reducing the rate of infection transmission in the population.



FIGURE 2: Effects of various intervention strategies on the control reproduction number.



FIGURE 3: Effects of varying the screening rate for latent infected individuals on the control reproduction number.

4.3. Effects of Asymptomatic Infectious Screening on the Symptomatic Population. Figure 6 illustrates the effects of varying the screening rate for asymptomatic infectious individuals on the symptomatic population. It is observed that an increase in screening for asymptomatic infectious individuals significantly reduces the symptomatic population. This reduction is attributed to the decreased progression of the asymptomatic infectious population to severe pulmonary tuberculosis disease. Asymptomatic infectious individuals often delay seeking healthcare and are not promptly identified for pulmonary tuberculosis testing. However, their detection is crucial for diagnosing the less advanced form of pulmonary tuberculosis and facilitating early treatment. Ultimately, this contributes to the reduction of disease transmission, decreased case fatality, and prevention of adverse consequences of the disease.

4.4. Effects of Screening Latent Infected on the Asymptomatic Infectious Population. Figure 7 illustrates the effects of screening latent infected individuals on the asymptomatic



FIGURE 4: Effects of varying the screening rate for asymptomatic infectious individuals on the control reproduction number.



FIGURE 5: Effects of varying the treatment rate for symptomatic infectious individuals on the control reproduction number.

infectious population. It is observed that an increase in screening of latent infected individuals results in a decrease in the asymptomatic infectious population. This reduction is due to decreased reactivation of latent infections as a result of treatment. The decrease in the asymptomatic infectious population reduces the incessant spread of infections to susceptible populations. Therefore, screening the latent infected population proves to be an effective strategy for reducing the transmission of infections in the population.



FIGURE 6: Effects of varying the screening rate for asymptomatic individuals on the symptomatic infectious population.



FIGURE 7: Effects of varying the screening rate for latent infected individuals on the asymptomatic infectious population.

#### 5. Conclusion

In this paper, a mathematical model of pulmonary tuberculosis incorporating an asymptomatic infectious population has been formulated. The asymptomatic infectious population was targeted since they spread infections incessantly to susceptible populations without being noticed, thus contributing to a high rate of infection transmission. This was based on research findings from a survey conducted in Kenya in 2016, which showed that 26% of active pulmonary tuberculosis cases were asymptomatic infectious and did not seek medical care. The model considered intervention strategies that include vaccination, screening of both latent infected and asymptomatic infectious populations, and treatment of all forms of pulmonary tuberculosis disease. Qualitative as well as numerical analyses have been performed to determine effective intervention strategies that reduce the transmission of pulmonary tuberculosis infections in the population. However, the limitations of this model, which will be considered in our next research paper, include the impact of natural immunity on the progression of individuals from latent infection to pulmonary tuberculosis disease and the waning of vaccine efficacy.

Qualitative and numerical results demonstrate that increasing the screening of asymptomatic and latently infected individuals reduces the transmission of infections to the susceptible population. The numerical analysis indicates that the combination of vaccination, screening, and treatment of all forms of pulmonary tuberculosis disease is the most effective intervention in decreasing disease transmission. Furthermore, the results suggest that a combination of screening and treatment of all forms of pulmonary tuberculosis disease is more effective than a combination of vaccination and treatment of symptomatic infectious individuals alone. Treating the symptomatic population alone is identified as the least effective intervention for curtailing infection transmission in the susceptible population.

Therefore, this study recommends that more attention should be directed toward screening and treatment of latent infected and the asymptomatic infectious populations. Screening and treating latent infections reduce the development of pulmonary tuberculosis disease and consequently decreases the rate of infection transmission in the population. Additionally, screening and treating the asymptomatic infectious population reduce the incessant spread of infections to susceptible individuals and, consequently, decreases the rate of infection transmission.

The limitations of this study, which could be considered in future studies, include the optimal control theory in the presence of vaccination, screening, and treatment, as well as the stochastic- and fractional-order approaches of the model.

#### **Data Availability**

The data used to support the findings of this study are included in the article.

#### **Conflicts of Interest**

The authors declare that there are no conflicts of interest regarding publication of this paper.

#### References

- H. W. Hethcote, "The mathematics of infectious diseases," Society for Industrial and Applied Mathematics, vol. 42, no. 4, pp. 599–653, 2000.
- [2] N. C. Grassly and C. Fraser, "Mathematical models of infectious disease transmission," *Nature Reviews Microbiology*, vol. 6, no. 6, pp. 477–487, 2008.
- [3] P. Yan and G. Chowell, Quantitative Methods for Investigating Infectious Disease Outbreaks, Springer, Cham, Switzerland, 2019.
- [4] World Health Organization, *Global Tuberculosis Report 2019*, World Health Organization, Geneva, Switzerland, 2020.
- [5] R. M. Anderson and R. M. May, *Infectious Diseases of Humans: Dynamics and Control*, Oxford University Press, Oxford, UK, 1991.
- [6] O. Diekmann, H. Heesterbeek, and T. Britton, Mathematical Tools for Understanding Infectious Disease Dynamics, Princeton University Press, Princeton, NJ, USA, 2013.
- [7] M. Martcheva, An Introduction to Mathematical Epidemiology, Springer, New York, NY, USA, 2015.
- [8] E. Vynnycky and R. White, An Introduction to Infectious Disease Modelling, OUP oxford, Oxford, UK, 2010.
- [9] W. O. Kermack and A. G. McKendrick, "A contribution to the mathematical theory of epidemics," *Proceedings of the Royal Society of London-Series A: Containing Papers of a Mathematical and Physical Character*, vol. 115, no. 772, pp. 700–721, 1927.
- [10] C. E. Barry, H. I. Boshoff, V. Dartois et al., "The spectrum of latent tuberculosis: rethinking the biology and intervention strategies," *Nature Reviews Microbiology*, vol. 7, no. 12, pp. 845–855, 2009.
- [11] World Health Organization, *Global Tuberculosis Report 2022*, World Health Organization, Geneva, Switzerland, 2023.
- [12] Knbs, "Statistical abstracts," 2020, https://www.knbs.or.ke/ wpdmpro=statistical-abstract-2020.
- [13] Unep, "Sustainable development Goals and the 2030 agenda: why environmental sustainability and gender equality are so important to reducing poverty and inequalities-UNEP perspectives issue No. 17," 2015, https://wedocs.unep.org/20.500. 11822/7464.
- [14] J. Andrawus, F. Eguda, I. G. Usman et al., "A mathematical model of a tuberculosis transmission dynamics incorporating first and second line treatment," *Journal of Applied Sciences & Environmental Management*, vol. 24, no. 5, pp. 917–922, 2020.
- [15] J. M. Trauer, J. T. Denholm, and E. S. McBryde, "Construction of a mathematical model for tuberculosis transmission in highly endemic regions of the Asia-Pacific," *Journal of Theoretical Biology*, vol. 358, pp. 74–84, 2014.
- [16] D. Boccia, W. Rudgard, S. Shrestha et al., "Modelling the impact of social protection on tuberculosis: the S-PROTECT project," *BMC Public Health*, vol. 18, no. 1, p. 786, 2018.
- [17] S. Athithan and M. Ghosh, "Mathematical modelling of TB with the effects of case detection and treatment," *International Journal of Dynamics and Control*, vol. 1, no. 3, pp. 223–230, 2013.
- [18] E. E. Salpeter and S. R. Salpeter, "Mathematical model for the epidemiology of tuberculosis, with estimates of the reproductive number and infection-delay function," *American Journal of Epidemiology*, vol. 147, no. 4, pp. 398–406, 1998.

- [19] R. M. Houben and P. J. Dodd, "The global burden of latent tuberculosis infection: a re-estimation using mathematical modelling," *PLoS Medicine*, vol. 13, no. 10, p. e1002152, 2016.
- [20] L. N. Nkamba, T. T. Manga, F. Agouanet, and M. L. Mann Manyombe, "Mathematical model to assess vaccination and effective contact rate impact in the spread of tuberculosis," *Journal of Biological Dynamics*, vol. 13, no. 1, pp. 26–42, 2019.
- [21] F. Sulayman, F. A. Abdullah, and M. H. Mohd, "An SVEIRE model of tuberculosis to assess the effect of an imperfect vaccine and other exogenous factors," *Mathematics*, vol. 9, no. 4, p. 327, 2021.
- [22] J. P. Aparicio and C. Castillo Chavez, "Mathematical modelling of tuberculosis epidemics," *Mathematical Biosciences* and Engineering: MBE, vol. 6, no. 2, pp. 209–237, 2009.
- [23] S. Kasereka Kabunga, E. F. Doungmo Goufo, and V. Ho Tuong, "Analysis and simulation of a mathematical model of tuberculosis transmission in Democratic Republic of the Congo," Advances in Difference Equations, vol. 2020, no. 1, p. 642, 2020.
- [24] Government of Kenya, Kenya Tuberculosis Prevalence Survey, Ministry of Health, Kenya, 2016.
- [25] P. Van den Driessche and J. Watmough, "Reproduction numbers and sub-threshold endemic equilibria for compartmental models of disease transmission," *Mathematical Biosciences*, vol. 180, no. 1-2, pp. 29–48, 2002.
- [26] M. A. Safi and S. M. Garba, "Global stability analysis of SEIR model with Holling type II incidence function," *Computational and Mathematical Methods in Medicine*, vol. 2012, Article ID 826052, 8 pages, 2012.
- [27] P. Van den Driessche, "Reproduction numbers of infectious disease models," *Infectious disease modelling*, vol. 2, no. 3, pp. 288–303, 2017.
- [28] J. P. La Salle, "The stability of dynamical systems," Society for Industrial and Applied Mathematics, SIAM, Philadelphia, PA, USA, 1976.
- [29] O. D. Makinde and K. O. Okosun, "Impact of chemo-therapy on optimal control of malaria disease with infected immigrants," *Biosystems*, vol. 104, no. 1, pp. 32–41, 2011.
- [30] National Tuberculosis Leprosy and Lung Disease Program, Annual Report, Ministry of Health, Kenya, 2021.
- [31] Knbs, Statistical Abstract 2023, KNBS, Nairobi, Kenya, 2023.
- [32] Knbs and Icf, Kenya Demographic and Health Survey 2022. Key Indicators Report, KNBS and ICF, Nairobi, Kenya, and Rockville, MY, USA, 2023.
- [33] National Tuberculosis Leprosy and Lung Disease Program, National Strategic Plan for Tuberculosis, Leprosy and Lung Health 2019-2023, Ministry of Health, Kenya.
- [34] National Tuberculosis Leprosy and Lung Disease Program, *Annual Report*, Ministry of Health, Kenya, 2022.
- [35] M. Enos, J. Sitienei, J. Ong'ang'o et al., "Kenya tuberculosis prevalence survey 2016: challenges and opportunities of ending TB in Kenya," *PLoS One*, vol. 13, no. 12, p. e0209098, 2018.
- [36] National Tuberculosis Leprosy and Lung Disease Program, Kenya Latent Tuberculosis Infection Policy 2020, Ministry of health, Kenya.
- [37] G. Chowell and J. M. Hyman, Mathematical and Statistical Modeling for Emerging and Re-emerging Infectious Diseases, Springer, Cham, 2016.