




Impact of saturated treatments on HIV-TB dual epidemic as a consequence of COVID-19: optimal control with awareness and treatment

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Abstract In this study, we propose an HIV-TB co-infection model by considering the treatment provision limitation induced by recent COVID-19 pandemic that impacts this dual epidemic immensely, assimilating the significance of educational attempts. We analyze the model and its submodels with single infections individually. We obtain the awareness-induced basic reproduction numbers and discuss the global stability of disease-free equilibrium when provision limitation is zero. We observe that the submodels exhibit forward as well as backward bifurcations under provision restriction. Further, we derive thresholds for resource limitations regulating the dynamical behavior of the systems while analyzing the stability of endemic equilibrium of the models with single infections. Sophisticated simulation approaches are implemented to discover the influences of provision-restricted medication and awareness on dual epidemic. Our findings convey the persistence of co-infection though the basic reproduction number is below unity, if the provision restric-

tion remains uncurbed. An observable insight is that, in spite of having epidemic threshold less than unity and no limitation in TB treatment, co-infection relapses and persists in the population, when there is no awareness attempt. Numerical findings emphasize the urgent need of increased treatment accessibility and importance of awareness in the current situation. Moreover, an optimization problem incorporating treatment and awareness controls is formulated and solved to find the ideal strategy to manage HIV-TB co-epidemic that recommends to diminish the medical resource limitation to get the enormous impact in dominating the adversity caused by COVID-19.

Keywords HIV-TB dual epidemic · Provision limitation · Educational attempt · Awareness-induced basic reproduction number · Optimal control · COVID-19

1 Introduction

Acquired immunodeficiency syndrome (AIDS) is a chronic assuredly intimidating condition caused by human immunodeficiency virus (HIV). HIV enfeeble human immune system by attacking and destroying CD4 cells that creates difficulty for the body to fight off infectious diseases and makes a person more vulnerable to other infections. If left untreated, HIV gradually demolishes the immunity structure and makes progress to AIDS. It can be transmitted only through

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contact with HIV-infected body fluids during unprotected sex or by sharing injection drug equipment or via HIV-contaminated blood transfusion or while pregnancy. Antiretroviral therapy (ART) is used to treat HIV infection, that protects the immune system and intercepts HIV infection from advancing to AIDS and reduces the risk of HIV transmission, as infected people with undetectable viral load (< 50 copies/ml blood) have effectively no probability of transferring HIV. Around the globe, an estimated 37.7 million people are infected with HIV [1].

TB (Tuberculosis) is a terrifying infectious bacterial disease, caused by mycobacterium tuberculosis. It is an airborne disease, as the bacteria spread by airborne respiratory droplets of an infectious individual while coughing or sneezing or can also be transmitted by saliva [2]. There are two conditions of TB infection, latent TB and active TB. In latent condition, TB bacteria exist within the body in a less quantity, so the immune system remains under control and do not cause any symptom. If latent TB endures untreated, it becomes fatal TB disease which is highly infectious. TB is curable with antimicrobial drugs [3]. Globally, an estimated 7.1 million people were reported to have been diagnosed in 2019, 9.96 million people got infected with TB in 2020, and 63 million lives were saved through TB diagnosis and treatment during 2000–2019 [4].

TB is synergistically related to HIV/AIDS where each augment the dreadfulness of other, and it can assuredly be a murderous combination which is capable of causing deaths if remains untreated. HIV-infected people are 30 times more presumably to get TB infection than a typically healthy person. A people infected with latent TB is able to develop active TB at least 10 times more in the presence of HIV. TB disease is more vulnerable to HIV-infected individuals because of their weakened immune system. An HIV-TB co-infection can accelerate the progression from HIV to AIDS. In 2020, an estimated 12.6 million people are co-infected globally [1,3]. The medications for HIV and TB combinedly can reduce the menace of co-infection. An HIV-infected person should continue ART in spite of being diagnosed with TB at the same time, but a specific attention should be given to avoid the potential drug interaction between ART and TB drugs. Isoniazid preventive therapy (IPT) is the most impactful medical support to HIV patients for preventing the progression from latent to active TB and reinfection. Around

4.1 million people are provided TB preventive treatment worldwide in 2019; among them 3.5 million are surviving with HIV. But, in developing countries like India and South Africa, the recent rates of getting TB preventive medications are 25 and 18%, respectively, which are not much satisfactory [4]. Moreover, some evidences show that ART can also prevent HIV-infected humans from acquiring TB disease by 70–90% [5]. In case of co-infection, it is necessary to start implementing ART as early as possible without detaining it until the completion of TB treatment as this strategy causes higher mortality rate [6]. Patients with HIV-TB co-infection acquire 1.8 times higher probability of death than a singly infected person [7]. HIV-TB combinedly impose an enormous burden on public health-care system and place specific diagnostic and medicinal challenges, especially in countries facing scarcity of resources. “End TB Strategy” by WHO targeted 20% reduction in TB incidence rate and 35% decrement in number of TB-induced deaths. But, it came across as only 9 and 14%, respectively, in which lack of awareness and treatment provision limitation contribute a lot. Nearly 14 million humans are victims of this dual epidemic worldwide and 86–90% of them are dense in high TB-HIV-loaded countries [8].

HIV-TB dual pandemic is exercised by numerous mathematical models [9–12]. Gakkhar and Chavda [13] showed that coexistence of HIV-TB cannot remain longer in population in spite of having basic reproduction number greater than unity, single disease can persist only. Agosto and Adekunle [14] studied an optimal control model considering two strains of TB (drug-sensitive and drug-resistant) to examine the effect of different control strategies. They showed that combined strategy including prevention of treatment failure in case of drug-sensitive TB and treatment of drug-resistant TB is the most effective one. Mallela et al. [15] analyzed a model to examine early and late treatment of HIV during TB treatment in case of co-infection, impacting new infection, HIV-related deaths and IRIS cases. Awoke and Kassa [16] formulated an optimal control model by taking into account prevalence depending behavior change incorporating treatment. They showed that combination of prevention and treatment is the best impactful optimal strategy. There are also some mathematical models on HIV-TB co-infection involving resource limitation matter. It can be noted that in all the aforementioned models, the treatment rate has been taken as constant following

Holling type-I function. In [17], authors have investigated an SIR model, where they have taken the treatment rate as Holling type-II functional response. Dubey et al. [18] considered Holling type-III and type-IV treatment rates in an SEIR model. In [19], authors have studied an SIR model by introducing Beddington–DeAngelis-type function as incidence rate together with Holling type-II treatment rate. Various modeling approaches have been introduced in some recent research papers [20–22].

COVID-19 pandemic has impacted the socio-economic structure badly; one of its unavoidable circumstances is disruption in medical resources. Noticeable retardation is found in TB medication encompassing preliminary care to hospitalization due to limited provision in health workers, medicines, pharmaceutical structure, finances, etc. Lockdown and public health protocols of COVID-19 situation implement tough challenges on conventional TB management. This restrictions can lead to a 13% increase of TB deaths [23]. According to WHO's survey, it has been directed onwards in International AIDS Society's (IAS) biannual conference that 73 countries are facing huge crisis because of the unavailability of antiretroviral medicines due to COVID-19 pandemic and 24 countries are identified as having an immensely feeble stock or upsetting supply of life-saving ARV drugs. A modeling investigation by WHO and UNAIDS predicted that a half-year disturbance in HIV medication service could double AIDS-induced mortality in Sub-Saharan Africa. Close-down of transportation hampers the suppliers of ARVs to reach the destination and COVID-19 pandemic situation causes limited access to medical service. Thus, the most needy people are unable to get prevention and testing facilities. Due to this, global target to suppress HIV is impeded as the estimated yearly new infections has been fastened at 1.7 million since last two years. There seems low-key reduction in HIV deaths from 7,30,000 in 2018 to 6,90,000 in 2019 [24]. Apparently, HIV-TB dual epidemic is directly impacted by COVID-19 situation. Thus, it is highly essential to take into consideration the provision limitation of HIV and TB treatment. That is why, we introduce saturated treatment rates in our model following Holling type-II functional response. Moreover, awareness plays a vital role in managing this dual epidemic by limiting risky behavior of susceptible individuals, especially in case of provision restriction situation. Almost 19% HIV-infected persons don't even know their condition [25].

It is very unfortunate that a lot of people surviving with HIV lead to TB related death and a huge portion of them don't even know their status or are not enough aware to start proper co-infection treatment within right time [26]. This necessitates enough awareness attempt to educate people about HIV, TB and most importantly to make them aware of preventive mechanism, diagnostic and pharmaceutical interventions to manage HIV-TB co-infection.

2 The model formulation

The main purpose of our present investigation is to advert the disruption of HIV-TB service ascribed to COVID-19. For this purpose, a fifteen-dimensional deterministic HIV-TB co-infection model has been formulated absorbing nonlinear treatment rate and awareness. At any time $t > 0$, let the total number of population per unit area in a considered domain be $N(t)$. The whole population is classified into fifteen mutually disjoint divisions (including four susceptible cohorts) viz. susceptible human population uneducated about HIV/AIDS and TB both, S_U^{TH} ; susceptible human population educated about HIV/AIDS only, S_E^H ; susceptible humans educated about TB only, S_E^T ; susceptible humans educated about HIV/AIDS and TB both, S_E^{TH} ; HIV-infected human population, H_1 ; AIDS-infected human population, H_2 ; latent TB-infected human population, L ; active TB-infected human population, I ; HIV-infected human population who are coinfecting with latent TB, L_{H_1} ; AIDS-infected human population who are coinfecting with latent TB, L_{H_2} ; active TB-infected human population who are coinfecting with HIV, I_{H_1} ; active TB-infected human population who are coinfecting with AIDS, I_{H_2} ; human population treated for HIV/AIDS only, T_H ; human population treated for TB only, T_T ; human population treated for HIV/AIDS and TB both, T_{TH} . We formulate our HIV-TB mathematical model based on the following assumptions.

1. Uneducated susceptible class has recruitment at a constant rate Λ , and the recruited individuals are completely unaware of these two diseases.
2. Uneducated susceptible individuals are consuming HIV education at a rate $\alpha_E^H = \frac{\alpha_1 R_1}{1+h_1 R_1}$, TB education at a rate $\alpha_E^T = \frac{\alpha_2 R_2}{1+h_2 R_2}$ and education about HIV-TB co-infection at a rate $\alpha_E^{TH} =$

$\frac{\alpha_3 \left(\frac{R_1+R_2+R_3}{3} \right)}{1+h_3 \left(\frac{R_1+R_2+R_3}{3} \right)}$, thereby moving to S_U^H , S_U^T and S_U^{TH} cohorts, respectively. HIV-educated susceptible humans can move to S_U^{TH} class by consuming TB education and similarly TB-educated individuals are capable to progress in S_U^{TH} class by consuming HIV education. Here, α_1 , α_2 and α_3 are information propagation rates for HIV, TB and HIV-TB co-infection, respectively; R_1 , R_2 and R_3 are the densities of educational attempts; and h_1 , h_2 and h_3 represent time-consuming parameters in case of awareness about HIV, TB and co-infection, accordingly.

3. Susceptible who are completely uneducated about HIV (i.e., S_U^H and S_U^T), acquire HIV infection by unmediated contact with HIV/AIDS-infected population including co-infected individuals at an effective infection rate,

$$\beta_H = \frac{\lambda_H[(H_1 + L_{TH_1} + I_{TH_1}) + \eta_1(H_2 + L_{TH_2} + I_{TH_2}) + \eta_2(T_H + T_{TH})]}{N}.$$

Since AIDS-infected individuals (including co-infection) are more vulnerable than HIV-infected persons (including co-infection) and infected people treated for HIV are less infectious than HIV infectives comparatively. Thus, we introduce two modification parameters $\eta_1 > 1$ and $\eta_2 < 1$, accordingly.

4. Susceptible humans who are unable to consume TB education (i.e., S_U^H and S_U^T) become infected from TB infectives including HIV-TB co-infected individuals at a rate of transmission

$$\beta_T = \frac{\lambda_T(I + I_{TH_1} + I_{TH_2})}{N}.$$

5. Information related to disease aspect changes exposed attitude of humans by adopting non-pharmacological intervention to avert infection. Educated susceptible individuals having knowledge about HIV/AIDS (i.e., S_E^H and S_E^{TH}) will be afflicted with HIV at a reduced rate $\beta_1\beta_H$, where β_1 is expressed as the measure of education impact depending on the decreased probability of risky human practice and efficacy of education, i.e., $\beta_1 = c_1(1 - \epsilon)$. TB disease transmits at a lesser rate $\beta_2\beta_H$, among susceptible humans having TB education (i.e., S_E^T and S_E^{TH}), where $\beta_2 = c_2(1 - \epsilon)$ with decreased probability of risky human practice

c_2 toward TB. Here, $(1 - \epsilon)$ represents the failure of educational movements.

6. People with HIV infection can develop AIDS without accessing any medical treatment at a rate δ and individuals infected with latent TB can progress to active TB infection at a rate ψ , if latent TB is evacuated to be oppressive.
7. HIV/AIDS is such a lethal disease that increases risk of other contagious diseases like TB by weakening the immune structure of body. As TB is an opportunistic infection, so HIV/AIDS-infected individuals become co-infected with TB at an increased rate $\phi\beta_T$. Similarly, TB infection provokes HIV transmission that induces an increased rate $\theta\beta_H$ for TB-infected persons to acquire HIV infection combinedly, where $\phi, \theta > 1$ are modification parameters.
8. HIV-TB is a deadly combination that can place someone's life in peril by accelerating each other's

progression in a damaged immune environment of the body. Therefore, HIV-infected individuals who acquire co-infection of TB disease escalate into AIDS at a rate $\sigma > \delta$, and co-infected people with latent TB and HIV/AIDS advance to active TB by a rapid activation at an increased rate $\zeta > \psi$.

9. Treatment rates for both HIV and TB are assumed to follow Holling type-II function considering limitation in the medical resources. HIV- and AIDS-infected people are accessing ART at a rate $\tau_{H_1} = \frac{lH_1}{1+v_1H_1}$ and $\tau_{H_2} = \frac{lH_2}{1+v_1H_2}$ ($l \geq 0$ is the initial treatment rate for HIV/AIDS), respectively. Similarly, people with latent TB and active TB are getting treatment at rates $\tau_L = \frac{mL}{1+v_2L}$ and $\tau_I = \frac{mI}{1+v_2I}$, respectively ($m \geq 0$ is the initial treatment rate for TB). People co-infected with HIV and TB are assumed to get medical intervention for co-infection at a rate $\tau_{H_T} = \frac{rH_T}{1+v_3H_T}$ (H_T represents co-infected classes, i.e., L_{H_1} , L_{H_2} , I_{H_1} and I_{H_2}) with initial treatment rate of co-infection $r \geq 0$. Here, $\frac{1}{1+v_1H_1}$ and $\frac{1}{1+v_1H_2}$ indicate the effects of hindrance for HIV and AIDS treatment, accordingly. The impact of restriction in latent TB treatment is indicated by $\frac{1}{1+v_2L}$ and active TB treat-

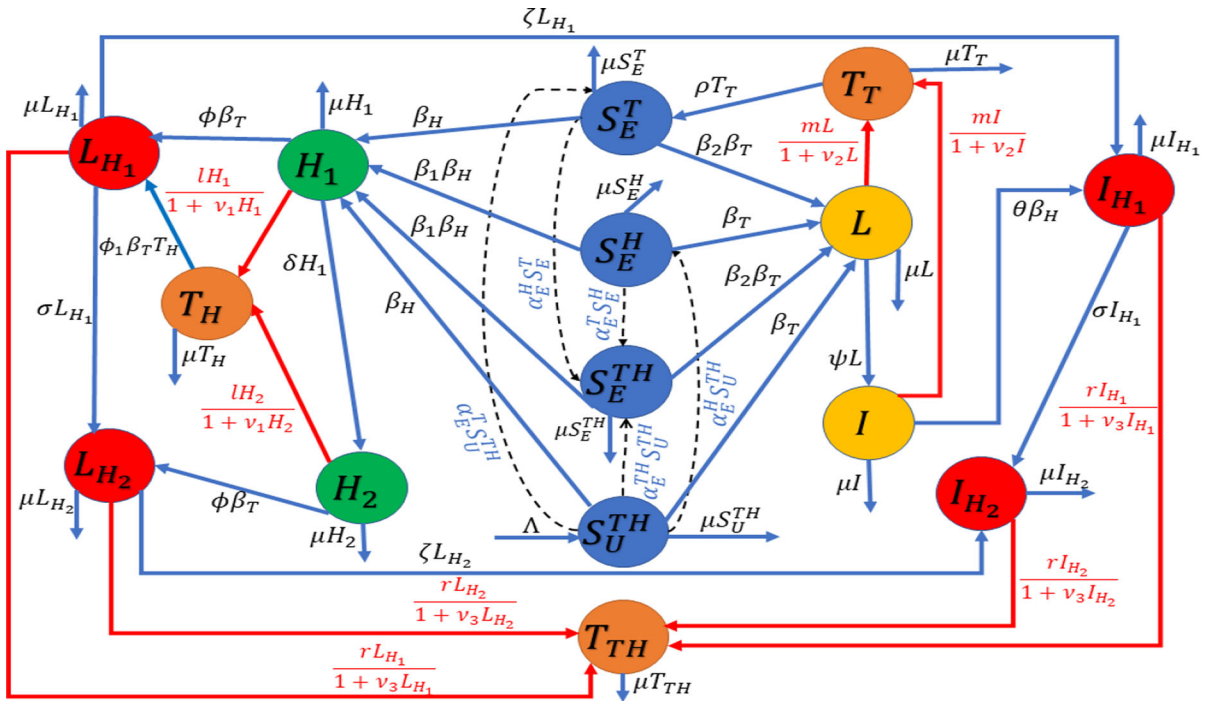


Fig. 1 Schematic diagram of the system (1) representing the impact of saturated treatments and awareness on HIV-TB dual epidemic. Here, the terms in red color represent the nonlinear treatment rates

ment by $\frac{1}{1+v_2I}$. Inhibition impact of co-infection treatment is represented by the term $\frac{1}{1+v_3I_H}$. Here, v_1, v_2 and v_3 are provision limitation parameters for HIV, TB and co-infection treatments, respectively. Clearly, the treatment rate is an increasing function of infected individuals with continuous differentiability, satisfying following conditions for an infected class I : (i) $\tau_0 = 0$, (ii) $\frac{d\tau_I}{dt} > 0$, $\frac{d^2\tau_I}{dt^2} < 0$, (iii) $\lim_{I \rightarrow \infty} \tau_I = \text{maximum treatment size}$.

10. Individuals are transferred to S_E^T class at a rate ρ from T_T class by losing their immunity with passage of time. This indicates the possibility of re-infection of individuals recovered from the TB disease. Moreover, individuals treated for HIV/AIDS can be infected with TB at a rate $\phi_1\beta_T$, where $1 < \phi_1 < \phi$ (since ART and IPT reduce the risk of getting TB infection while developing co-infection in comparison with untreated HIV/AIDS-infected people).
11. Per-capita natural mortality rate is μ in each cohort whereas due to the graveness of diseases there are extra mortalities for the individuals infected with

AIDS, TB and HIV-TB co-infection at rates d_H, d_T and d_{TH} , respectively.

In view of the above assumptions, a schematic diagram is depicted in Fig. 1, and we have the following system of differential equations for the HIV-TB coinfection:

$$\begin{aligned} \frac{dS_U^{TH}}{dt} &= \Lambda - (\alpha_E^H + \alpha_E^T + \alpha_E^{TH} + \beta_T + \beta_H + \mu)S_U^{TH}, \\ \frac{dS_E^H}{dt} &= \alpha_E^H S_U^{TH} - (\beta_1\beta_H + \beta_T + \alpha_E^T + \mu)S_E^H, \\ \frac{dS_E^T}{dt} &= \alpha_E^T S_U^{TH} - (\beta_2\beta_T + \beta_H + \alpha_E^H + \mu)S_E^T + \rho T_T, \\ \frac{dS_E^{TH}}{dt} &= \alpha_E^{TH} S_U^{TH} + \alpha_E^T S_E^H + \alpha_E^H S_E^T \\ &\quad - (\beta_1\beta_H + \beta_2\beta_T + \mu)S_E^{TH}, \\ \frac{dH_1}{dt} &= \beta_H S_U^{TH} + \beta_1\beta_H S_E^H + \beta_H S_E^T \\ &\quad + \beta_1\beta_H S_E^{TH} + \beta_H T_T - \phi\beta_T H_1 \\ &\quad - (\delta + \mu)H_1 - \tau_{H_1}, \\ \frac{dH_2}{dt} &= \delta H_1 - (\phi\beta_T + \mu + d_H)H_2 - \tau_{H_2}, \\ \frac{dL}{dt} &= \beta_T S_U^{TH} + \beta_T S_E^H + \beta_2\beta_T S_E^T \end{aligned}$$

$$\begin{aligned}
 & +\beta_2\beta_T S_E^{TH} - (\theta\beta_H + \psi + \mu)L - \tau_L, \\
 \frac{dI}{dt} & = \psi L - (\theta\beta_H + \mu + d_T)I - \tau_I, \\
 \frac{dL_{H_1}}{dt} & = \theta\beta_H L + \phi\beta_T H_1 \\
 & + \phi_1\beta_T T_H - (\sigma + \mu + \zeta)L_{H_1} - \tau_{L_{H_1}}, \\
 \frac{dL_{H_2}}{dt} & = \phi\beta_T H_2 + \sigma L_{H_1} \\
 & - (\zeta + d_H + \mu)L_{H_2} - \tau_{L_{H_2}}, \\
 \frac{dI_{H_1}}{dt} & = \theta\beta_H I + \zeta L_{H_1} \\
 & - (\sigma + d_T + \mu)I_{H_1} - \tau_{I_{H_1}}, \\
 \frac{dI_{H_2}}{dt} & = \sigma I_{H_1} + \zeta L_{H_2} - (d_{TH} + \mu)I_{H_2} - \tau_{I_{H_2}}, \\
 \frac{dT_T}{dt} & = \tau_L + \tau_I - \beta_H T_T - (\mu + \rho)T_T, \\
 \frac{dT_H}{dt} & = \tau_{H_1} + \tau_{H_2} - \phi_1\beta_T T_H - \mu T_H, \\
 \frac{dT_{TH}}{dt} & = \tau_{L_{H_1}} + \tau_{L_{H_2}} + \tau_{I_{H_1}} + \tau_{I_{H_2}} - \mu T_{TH}. \tag{1}
 \end{aligned}$$

Here,

$$\begin{aligned}
 \tau_{H_1} & = \frac{lH_1}{1 + v_1 H_1}, \quad \tau_{H_2} = \frac{lH_2}{1 + v_1 H_2}, \\
 \tau_L & = \frac{mL}{1 + v_2 L}, \quad \tau_I = \frac{mT}{1 + v_2 I}, \\
 \tau_{L_{H_1}} & = \frac{rL_{H_1}}{1 + v_3 L_{H_1}}, \quad \tau_{L_{H_2}} = \frac{rL_{H_2}}{1 + v_3 L_{H_2}}, \\
 \tau_{I_{H_1}} & = \frac{rI_{H_1}}{1 + v_3 I_{H_1}}, \quad \tau_{I_{H_2}} = \frac{rI_{H_2}}{1 + v_3 I_{H_2}}.
 \end{aligned}$$

All the variables and parameters related to the model (1) are assumed to be nonnegative as the system dynamic observes human population. System (1) is to be analyzed with nonnegative initial conditions. The epidemiological meanings of parameters describing model (1) are mentioned in Table 1.

Let us consider the biologically-reasonable domain:

$$\begin{aligned}
 \Omega & = \left\{ (S_U^{TH}, S_E^H, S_E^T, S_E^{TH}, H_1, H_2, L, I, L_{TH_1}, \right. \\
 & \left. L_{TH_2}, I_{H_1}, I_{TH_2}, T_T, T_H, T_{TH}) \in \mathbb{R}_+^{15} : N \leq \frac{\Lambda}{\mu} \right\}.
 \end{aligned}$$

The dynamics of total population $N(t)$ is given by adding all the equations of model (1) to get,

$$\frac{dN}{dt} = \Lambda - \mu N - d_H H_2 - d_T I - d_H L_{H_2} - d_T I_{H_1} - d_{TH} I_{H_2} \leq \Lambda - \mu N.$$

Applying standard comparison theorem [29] and using the initial conditions, we have $N(t) \leq \frac{\Lambda}{\mu} + e^{-\mu t} [N(0) - \frac{\Lambda}{\mu}]$. Particularly, $N(t) \leq \frac{\Lambda}{\mu}$ whenever $N(0) \leq \frac{\Lambda}{\mu}$.

Therefore, every solution of the system (1) initiating in Ω remains therein for all $t \geq 0$. Hence, Ω is a positive invariant and attracting region. It assures the epidemiological and mathematical well-posedness of the system (1) in Ω .

3 Dynamics of system (1) in the absence of TB

The HIV sub-model of the co-infection system (1) is obtained when human population is not infected with TB disease, i.e., when all the compartments of human population in system (1) other than S_U^H, S_E^H, H_1, H_2 and T_H are absent. The HIV sub-system, well-posed in a subdomain $\Omega_1 \subset \Omega$, is given as follows:

$$\begin{aligned}
 \frac{dS_U^H}{dt} & = \Lambda - (\alpha_E^H + \beta_H + \mu)S_U^H, \\
 \frac{dS_E^H}{dt} & = \alpha_E^H S_U^H - (\beta_1\beta_H + \mu)S_E^H, \\
 \frac{dH_1}{dt} & = \beta_H S_U^H + \beta_1\beta_H S_E^H - (\delta + \mu)H_1 - \tau_{H_1}, \\
 \frac{dH_2}{dt} & = \delta H_1 - (\mu + d_H)H_2 - \tau_{H_2}, \\
 \frac{dT_H}{dt} & = \tau_{H_1} + \tau_{H_2} - \mu T_H. \tag{2}
 \end{aligned}$$

Here, $\beta_H = \frac{\lambda_H(H_1 + \eta_1 H_2 + \eta_2 T)}{N}$.

3.1 Disease-free equilibrium (DFE) and its stability

The disease-free equilibrium of HIV-only model can be obtained by equating the right-hand side of system (2) to zero, when the disease does not exist in the population, and it is given by

$$\begin{aligned}
 \chi_0^H & = (\widehat{S}_U^H, \widehat{S}_E^H, \widehat{H}_1, \widehat{H}_2, \widehat{T}_H) \\
 & = \left(\frac{\Lambda}{\alpha_E^H + \mu}, \frac{\Lambda\alpha_E^H}{\mu(\alpha_E^H + \mu)}, 0, 0, 0 \right).
 \end{aligned}$$

Therefore, at DFE, we have $\widehat{N} = \frac{\Lambda}{\mu}$. Now, we derive the expression for basic reproduction number by next-generation matrix approach [30]. The matrices \mathcal{F} and \mathcal{V} associated with infection terms and rest of the left transfer terms, respectively, are computed as,

$$\mathcal{F}(x) = \begin{bmatrix} \beta_H S_U^H + \beta_1\beta_H S_E^H & & & \\ & 0 & & \\ & 0 & & \\ & 0 & & \\ & 0 & & \end{bmatrix},$$

Table 1 Epidemiological meanings of parameters in system (1) and their values used for numerical simulations

Parameters	Descriptions	Values	References
A	Rate of recruitment into the class of uneducated susceptible population	10353	[27]
α_1	Rate of propagating information about HIV	0.18	[16]
α_2	Rate of propagating information about TB	0.12	[16]
α_3	Average rate of propagating information about HIV and TB both	0.15-0.25	Assumed
R_1, R_2	Densities of awareness attempt about HIV and TB, respectively	0.5, 0.6	Variable
R_3	Density of awareness attempt about co-infection	0.32	Variable
h_1, h_2	Time-consuming parameters for HIV and TB education, respectively	0.657, 0.438	Assumed
h_3	Time-consuming parameter for education about HIV-TB co-infection	0.45-0.876	Assumed
ϵ	Efficacy of education	0.093	Assumed
λ_H	Effective transmission rate of HIV	0.35	Variable
λ_T	Effective transmission rate of TB	1.5	Variable
c_1	Reduced probability of risky human behavior after getting education about HIV	0.7	Assumed

Table 1 continued

Parameters	Descriptions	Values	References
c_2	Reduced probability of risky human behavior after getting education about TB	0.6	Assumed
δ	Rate of progression from HIV to AIDS	0.1	[16]
ψ	Rate of progression from latent TB to active TB	0.213	[28]
σ	Increased rate of progression from HIV to AIDS in case of coinfection	0.25	[28]
ζ	Increased rate of progression from latent TB to active TB in case of coinfection	0.25	[28]
l	Initial treatment rate of HIV/AIDS-infected individuals	0.62	Variable
m	Initial treatment rate for TB-infected individuals	0.7	[28]
r	Initial treatment rate for coinfecting individuals	0.66	Variable
ν_1	Provision limitation parameter for HIV/AIDS	0.06	Assumed
ν_2	Provision limitation parameter for TB	0.07	[28]
ν_3	Provision limitation parameter for co-infection	0.065	Assumed
ρ	Rate of transfer of individuals in the class of S_E^T from T_T	0.2	[28]
μ	Natural death rate of humans	0.02	[16]
d_H	HIV-induced death rate	0.3	[28]
d_T	TB-induced death rate	0.1	[28]
d_{TH}	HIV-TB coinfection-induced death rate	0.39996	[28]
$\phi, \phi_1, \theta, \eta_1, \eta_2$	Modification parameters	1.07, 1.007, 1.03, 1.08, 0.02	Assumed

$$\mathcal{V}(x) = \begin{bmatrix} (\delta + l + \mu)H_1 \\ (\mu + d_H)H_2 + \tau_{H_2} - \delta H_1 \\ \mu T_H - \tau_{H_1} - \tau_{H_2} \\ (\alpha_E^H + \beta_H + \mu)S_U^H - \Lambda \\ (\beta_1 \beta_H + \mu)S_E^H - \alpha_E^H S_U^H \end{bmatrix}.$$

The corresponding Jacobian matrices are obtained, respectively, as

$$F = \begin{bmatrix} \lambda_H \left(\frac{\mu + \beta_1 \alpha_E^H}{\mu + \alpha_E^H} \right) & \eta_1 \lambda_H \left(\frac{\mu + \beta_1 \alpha_E^H}{\mu + \alpha_E^H} \right) & \eta_2 \lambda_H \left(\frac{\mu + \beta_1 \alpha_E^H}{\mu + \alpha_E^H} \right) \\ 0 & 0 & 0 \\ 0 & 0 & 0 \end{bmatrix},$$

$$V = \begin{bmatrix} \delta + l + \mu & 0 & 0 \\ -\delta & l + \mu + d_H & 0 \\ -l & -l & \mu \end{bmatrix}.$$

Thus,

$$FV^{-1} = \begin{bmatrix} K_1 & K_2 & 0 \\ 0 & 0 & 0 \\ 0 & 0 & 0 \end{bmatrix}.$$

Here,

$$K_1 = \lambda_H \left[\frac{(\mu + \beta_1 \alpha_E^H)}{(\mu + \alpha_E^H)(\delta + l + \mu)} \right]$$

$$\left[1 + \frac{\delta \eta_1}{l + \mu + d_H} + \frac{\delta l \eta_2}{\mu(l + \mu + d_H)} + \frac{\eta_2 l}{\mu} \right]$$

and

$$K_2 = \lambda_H \left[\frac{(\mu + \beta_1 \alpha_E^H)}{(\mu + \alpha_E^H)(\delta + l + \mu)} \right] \left[\eta_1 + \frac{\eta_2}{\mu} \right].$$

Note that FV^{-1} produces the awareness-induced basic reproduction number for HIV-only model and is defined as

$$\begin{aligned} \mathcal{R}_E^H &= \bar{\rho}(FV^{-1}) \\ &= \frac{\lambda_H(\mu + \beta_1\alpha_E^H)[\mu(l + \mu + d_H) + \delta(\mu\eta_1 + l\eta_2) + l\eta_2(l + \mu + d)]}{\mu(\mu + \alpha_E^H)(\delta + l + \mu)(l + \mu + d_H)}. \end{aligned} \tag{3}$$

Here, $\bar{\rho}$ is the spectral radius of next-generation matrix. Moreover, in the absence of any awareness attempt, the basic reproduction number transforms to \mathcal{R}_0^H , which is given by

$$\mathcal{R}_0^H = \frac{\lambda_H[(\mu + l\eta_2)(l + \mu + d_H) + \delta(\mu\eta_1 + l\eta_2)]}{\mu(l + \mu + d_H)(\delta + l + \mu)}. \tag{4}$$

Following [30], we have the following theorem.

Theorem 1 *The disease-free equilibrium of HIV model (2) is locally asymptotically stable if $\mathcal{R}_E^H < 1$ and unstable if $\mathcal{R}_E^H > 1$.*

Regarding the global stability of disease-free equilibrium χ_0^H , we have the following theorem.

Theorem 2 *If the treatment provision of HIV/AIDS has no restriction ($v_1 = 0$), the disease-free equilibrium χ_0^H of HIV model (2) is globally asymptotically stable in Ω_1 for $\mathcal{R}_E^H \leq 1$ and $\mathcal{R}_0^H \leq 1$.*

For the proof of this theorem, see Appendix A.

3.2 Impact of public health education on \mathcal{R}_E^H

Let $E_1 = \frac{\widehat{S}_E^H}{N}$ be the proportion of individuals educated about HIV/AIDS in disease-free environment. From equation (3), we have

$$\mathcal{R}_E^H = \lambda_H [1 + (\beta_1 - 1)E_1] \left[\frac{\mu(l + \mu + d_H) + \delta(\mu\eta_1 + l\eta_2) + l\eta_2(l + \mu + d_H)}{\mu(\delta + l + \mu)(l + \mu + d_H)} \right].$$

After partially differentiating the above equation, we get

$$\begin{aligned} \frac{\partial \mathcal{R}_E^H}{\partial E_1} &= \lambda_H(\beta_1 - 1) \left[\frac{\mu(l + \mu + d_H) + \delta(\mu\eta_1 + l\eta_2) + l\eta_2(l + \mu + d_H)}{\mu(\delta + l + \mu)(l + \mu + d_H)} \right] \\ &= K(\beta_1 - 1), \end{aligned} \tag{5}$$

where

$$K = \lambda_H \left[\frac{\mu(l + \mu + d_H) + \delta(\mu\eta_1 + l\eta_2) + l\eta_2(l + \mu + d_H)}{\mu(\delta + l + \mu)(l + \mu + d_H)} \right] > 0.$$

As K is always positive, we have the following possibilities:

(1) $\frac{\partial \mathcal{R}_E^H}{\partial E_1} > 0$ if $\beta_1 > 1$; (2) $\frac{\partial \mathcal{R}_E^H}{\partial E_1} < 0$ if $\beta_1 < 1$;

(3) $\frac{\partial \mathcal{R}_E^H}{\partial E_1} = 0$ if $\beta_1 = 1$. Actually, β_1 is the measure of HIV/AIDS education impact on HIV transmission rate due to the change of hazardous human behavior toward HIV infection.

If the efficacy of health education is 100%, β_1 reaches the value zero as $\epsilon(100\%) = 1$ and $\beta_1 = c_1(1 - \epsilon)$, where c_1 is the reduced probability of risky human practice. As β_1 is always less than unity, $\frac{\partial \mathcal{R}_E^H}{\partial E_1} < 0$. This indicates that health education of HIV has negative impact on effective basic reproduction number \mathcal{R}_E^H .

Therefore, the influence of HIV education will be inimical only when the propagation of awareness is not perfect. Again, as $\beta_1 < 1$, from equations (3) and (4), we have $\frac{R_E^H}{R_0^H} = \frac{\mu + \beta_1\alpha_E^H}{\mu + \alpha_E^H} < \frac{\mu + \alpha_E^H}{\mu + \alpha_E^H} = 1$.

This indicates that proper HIV education always has positive impact in controlling the disease and hence reduces the burden of HIV/AIDS epidemic.

3.3 Local stability of endemic equilibrium and bifurcation analysis

The endemic equilibrium point of HIV sub-model (2) is denoted by $\chi_H^* = (S_U^{H*}, S_E^{H*}, H_1^*, H_2^*, T_H^*)$, and is obtained by setting the right-hand side of system (2) to zero, to get

$$\begin{aligned} S_U^{H*} &= \frac{\Lambda}{\alpha_E^H + \beta_H^* + \mu}, \\ S_E^{H*} &= \frac{\Lambda\alpha_E^H}{(\alpha_E^H + \beta_H^* + \mu)(\beta_1\beta_H^* + \mu)}, \\ H_1^* &= \frac{1}{\delta} \left[\frac{(l + \mu + d_H)H_2^* + v_1(\mu + d_H)H_2^{*2}}{1 + v_1H_2^*} \right], \end{aligned} \tag{6}$$

$$T_H^* = \frac{1}{\mu} \left[\frac{lH_1^*}{1 + v_1H_1^*} + \frac{lH_2^*}{1 + v_1H_2^*} \right]. \tag{7}$$

After adding all the equations of system (2), we get

$$\Lambda - \mu N^* - d_H H_2^* = 0. \tag{8}$$

At the endemic equilibrium point, we have

$$N^* \beta_H^* = \lambda_H (H_1^* + H_2^* + T_H^*). \tag{9}$$

We can find the value of H_2^* by substituting the values of S_U^* , S_E^* , H_1^* , H_2^* and T_H^* in Eq. (8). Now, from Eqs. (6–7), we can obtain H_1^* and T_H^* . Moreover, we can derive the force of infection β_H^* from Eq. (9). For $S_U^* > 0$, $S_E^* > 0$, $H_1^* > 0$, $H_2^* > 0$, $T_H^* > 0$, the feasibility of endemic equilibrium point χ_H^* is assured. Moreover, the equilibrium χ_H^* is unique if H_2^* and β_H^* are determined uniquely from Eqs. (8) and (9), respectively.

We have the following theorem regarding the local stability of endemic equilibrium point χ_H^* .

Theorem 3 *The unique endemic equilibrium point χ_H^* exists for $\mathcal{R}_E^H > 1$ and is locally asymptotically stable whenever $v_1 < \bar{v}_1$; the system (2) exhibits forward bifurcation with bifurcation parameter λ_H^* at $\mathcal{R}_E^H = 1$. Moreover, there exists a stable endemic equilibrium along with a stable disease-free equilibrium for $\mathcal{R}_E^H < 1$ whenever $v_1 > \bar{v}_1$ and system (2) exhibits a backward bifurcation, where \bar{v}_1 is defined in the proof.*

For the proof of this theorem, see Appendix B.

4 Dynamics of system (1) in the absence of HIV/AIDS

The TB sub-model of the co-infection system (1) is obtained when no HIV/AIDS is present in the human population, i.e., when all other classes are zero except S_U^T , S_E^T , L , I , T_T , and is given as follows:

$$\begin{aligned} \frac{dS_U^T}{dt} &= \Lambda - (\alpha_E^T + \beta_T + \mu)S_U^T, \\ \frac{dS_E^T}{dt} &= \alpha_E^T S_U^T - (\beta_2 \beta_T + \mu)S_E^T + \rho T_T, \\ \frac{dL}{dt} &= \beta_T S_U^T + \beta_2 \beta_T S_E^T - (\psi + \mu)L - \tau_L, \\ \frac{dI}{dt} &= \psi L - (\mu + d_T)I - \tau_I, \\ \frac{dT_T}{dt} &= \tau_L + \tau_I - (\mu + \rho)T_T. \end{aligned} \tag{10}$$

Here, $\beta_T = \lambda_T I/N$. System (10) is well-posed in the subdomain $\Omega_2 \subset \Omega$.

4.1 Disease-free equilibrium and its stability

The disease-free equilibrium point of the model (10) is obtained as

$$\begin{aligned} \chi_0^T &= (\widehat{S}_U^T, \widehat{S}_E^T, \widehat{L}, \widehat{I}, \widehat{T}_T) \\ &= \left(\frac{\Lambda}{\alpha_E^T + \mu}, \frac{\Lambda \alpha_E^T}{\mu(\alpha_E^T + \mu)}, 0, 0, 0 \right), \end{aligned}$$

which always exists in the system.

The awareness-induced basic reproduction number for model (10) is obtained, by following next-generation matrix approach [30], as

$$\begin{aligned} \mathcal{R}_E^T &= \bar{\rho}(FV^{-1}) \\ &= \frac{\lambda_T \psi (\mu + \beta_2 \alpha_E^T)}{(\mu + \alpha_E^T)(\psi + m + \mu)(m + \mu + d_T)}. \end{aligned} \tag{11}$$

It is apparent from the expression of \mathcal{R}_E^T that proper TB education always has a positive impact in curbing the disease and it reduces the burden of TB epidemic. In the absence of any educational attempt, \mathcal{R}_E^T takes the form

$$\mathcal{R}_0^T = \frac{\lambda_T \psi}{(\psi + m + \mu)(m + \mu + d_T)}. \tag{12}$$

Thus, we get the following result [30].

Theorem 4 *The disease-free equilibrium χ_0^T is locally asymptotically stable if $\mathcal{R}_E^T < 1$ and unstable if $\mathcal{R}_E^T > 1$.*

Regarding the global stability of χ_0^T , we have the following theorem.

Theorem 5 *The disease-free equilibrium χ_0^T is globally asymptotically stable (GAS) in Ω_2 for $\mathcal{R}_E^T \leq 1$ as well as for $\mathcal{R}_0^T \leq 1$ only when there is no limitation of TB treatment provision, i.e., $v_2 = 0$.*

Proof Let us consider the following Lyapunov function:

$$\mathcal{L}_2 = b_1 L + b_2 I,$$

where $b_1 = \psi$ and $b_2 = \psi + m + \mu$ with Lyapunov derivative,

$$\begin{aligned} \dot{\mathcal{L}}_2 &= b_1 \dot{L} + b_2 \dot{I} \\ &= b_1 [\beta_T S_U^T + \beta_2 \beta_T S_E^T - (\psi + \mu)L - \tau_L] \\ &\quad + b_2 [\psi L - (\mu + d_T)I - \tau_I] \\ &= \left[\frac{N \beta_T (\psi + m + \mu)(n + \mu + d_T)}{\lambda_T} \right] \end{aligned}$$

$$(\mathcal{R}_{eff}^T - 1), \quad (if \ v_2 = 0).$$

If TB treatment provision has no limitation, i.e., only when $v_2 = 0$, we must have $\dot{\mathcal{L}}_2 \leq 0$ for $\mathcal{R}_E^T \leq 1$.

Therefore, by LaSalle’s Invariance Principle [31], the disease-free equilibrium χ_0^T is globally asymptotically stable only when $v_2 = 0$. \square

4.2 Endemic equilibrium and its local stability

The endemic equilibrium of the model with TB only is $\chi_T^* = (S_U^{T*}, S_E^{T*}, L^*, I^*, T_T^*)$, whose components are obtained as follows:

$$S_U^{T*} = \frac{\Lambda}{\alpha_E^T + \beta_T + \mu}, \tag{13}$$

$$S_E^{T*} = \frac{\Lambda \alpha_E^T}{(\alpha_E^T + \beta_T + \mu)(\beta_2 \beta_T + \mu)} + \frac{\rho T_T^*}{\beta_2 \beta_T + \mu}, \tag{14}$$

$$L^* = \frac{1}{\psi} \left[\frac{(m + \mu + d_T)I^* + v_2(\mu + d_T)I^{*2}}{1 + v_2 I^*} \right], \tag{15}$$

$$T_T^* = \frac{1}{(\mu + \rho)} \left[\frac{mL^*}{1 + v_2 L^*} + \frac{mI^*}{1 + v_2 I^*} \right], \tag{16}$$

$$A - \mu N^* - d_T I^* = 0, \tag{17}$$

$$\beta_T^* = \frac{\lambda_T I^*}{S_U^{T*} + S_E^{T*} + L^* + I^* + T_T^*}. \tag{18}$$

The positive values of S_E^{T*} , L^* , I^* and T_T^* can be determined from Eqs. (13–16). Further, the force of TB infection β_T^* can be derived from Eq. (18). If the solution of equilibrium equations is unique, it refers to the unique existence of the endemic equilibrium χ_T^* .

Employing center manifold theory (Theorem 4.1 of [32]), we get the following result regarding the local stability of the endemic equilibrium χ_T^* .

Theorem 6 *The unique endemic equilibrium point χ_T^* exists for $\mathcal{R}_E^T > 1$ and is locally asymptotically stable whenever $v_2 < \bar{v}_2$, the system (10) undergoes a forward bifurcation at $\mathcal{R}_E^T = 1$ with bifurcation parameter λ_T^* . Further, there exists a stable endemic equilibrium and a stable disease-free equilibrium simultaneously for $\mathcal{R}_E^T < 1$ whenever $v_2 > \bar{v}_2$ and system (10) exhibits backward bifurcation, where*

$$\bar{v}_2 = \frac{\psi \mu \lambda_T^* (\psi + m + \mu) (\mu + \beta_2 \alpha_E^T)}{2m[\psi(\psi + m + \mu) + (m + \mu + d_T)^2] (\mu + \alpha_E^T)} \left[1 + \frac{\psi}{(m + \mu + d_T)} + \left(\frac{m}{m + \rho} \right) \left(1 + \frac{\psi}{m + \mu + d_T} \right) \right].$$

5 Analysis of HIV-TB co-infection system (1)

5.1 Disease-free equilibrium and its stability

Disease-free equilibrium of system (1) is obtained as,

$$\begin{aligned} \chi_0 &= (\widehat{S}_U^{TH}, \widehat{S}_E^H, \widehat{S}_E^T, \widehat{S}_E^{TH}, \widehat{H}_1, \widehat{H}_2, \widehat{L}, \widehat{I}, \widehat{L}_{H_1}, \\ &\quad \widehat{L}_{H_2}, \widehat{I}_{H_1}, \widehat{I}_{H_2}, \widehat{T}_T, \widehat{T}_H, \widehat{T}_{TH}) \\ &= \left(\frac{\Lambda}{(\alpha_E^H + \alpha_E^T + \alpha_E^{TH} + \mu)}, \right. \\ &\quad \frac{\Lambda \alpha_E^H}{(\alpha_E^T + \mu) (\alpha_E^H + \alpha_E^T + \alpha_E^{TH} + \mu)}, \\ &\quad \frac{\Lambda \alpha_E^T}{(\alpha_E^H + \mu) (\alpha_E^H + \alpha_E^T + \alpha_E^{TH} + \mu)}, \\ &\quad \frac{\Lambda \left(\alpha_E^{TH} + \frac{\alpha_E^H \alpha_E^T}{\alpha_E^H + \mu} + \frac{\alpha_E^H \alpha_E^T}{\alpha_E^T + \mu} \right)}{\mu (\alpha_E^H + \alpha_E^T + \alpha_E^{TH} + \mu)}, \\ &\quad \left. 0, 0, 0, 0, 0, 0, 0, 0, 0, 0 \right) \end{aligned}$$

The basic reproduction number of system (1) is derived by next-generation matrix method [30]. The awareness-induced basic reproduction number \mathcal{R}_E of the co-infection dynamics is obtained as,

$$\mathcal{R}_E = \bar{\rho}(FV^{-1}) = \max\{\mathcal{R}_E^H, \mathcal{R}_E^T\}. \tag{19}$$

If there is no educational attempt about HIV and TB, then \mathcal{R}_E is transformed into the following form:

$$\mathcal{R}_0 = \max\{\mathcal{R}_0^H, \mathcal{R}_0^T\}. \tag{20}$$

Thus, we get the following result [30].

Theorem 7 *The disease-free equilibrium point χ_0 is locally asymptotically stable if $\mathcal{R}_E < 1$ and unstable if $\mathcal{R}_E > 1$.*

5.1.1 Global stability of disease-free equilibrium χ_0

Following Castillo-Chavez et al. [33], we study global stability behavior of DFE χ_0 . System (1) can be rewritten as: $\frac{d\mathcal{K}}{dt} = G(\mathcal{K}, \mathcal{L})$, $\frac{d\mathcal{L}}{dt} = H(\mathcal{K}, \mathcal{L})$. Here, \mathcal{K} and \mathcal{L} represent the uninfected and infected individuals, respectively. For the global stability of equilibrium χ_0 , we must have $H(\mathcal{K}, \mathcal{L}) = A\mathcal{L} - \widehat{H}(\mathcal{K}, \mathcal{L})$, $\widehat{H}(\mathcal{K}, \mathcal{L}) \geq 0$ ($\mathcal{K}, \mathcal{L} \in \Omega$). Here, A represents an M-matrix.

For system (1), $\mathcal{L} = (H_1, H_2, L, I, L_{H_1}, L_{H_2}, I_{H_1}, I_{H_2}, T_H, T_{TH})^T \in \mathbb{R}^{10}$ and the matrix $\hat{H}(\mathcal{K}, \mathcal{L})$ is computed as

maximum rates of education consumptions. Maximum treatment rates are indicated as l_{max} , m_{max} and r_{max} ,

$$\hat{H}(\mathcal{K}, \mathcal{L}) = \begin{bmatrix} \lambda_H L_1 [(S_1 - \frac{S_U^{TH}}{N} - \beta_1 \frac{S_E^H}{N} - \frac{S_E^T}{N} - \beta_1 \frac{S_E^{TH}}{N})] + \phi \beta_T H_1 - \beta_H T_T - \frac{l v_1 H_1^2}{1+v_1 H_1} \\ \lambda_T L_2 [(S_1 - \frac{S_U^{TH}}{N} - \frac{S_E^H}{N} - \beta_2 \frac{S_E^T}{N} - \beta_2 \frac{S_E^{TH}}{N})] - \frac{m v_2 L^2}{1+v_2 L} \\ - \frac{l v_1 H_2^2}{1+v_1 H_2} \\ \theta \beta_H I - \frac{m v_2 I^2}{1+v_2 I} \\ -\theta \beta_H L - \phi \beta_T H_1 - \phi_1 \beta_T T_H - \frac{r v_3 L H_1^2}{1+v_3 L H_1} \\ -\phi \beta_T H_2 - \frac{r v_3 L H_2^2}{1+v_3 L H_2} \\ -\theta_H \beta_H I - \frac{r v_3 I H_1^2}{1+v_3 I H_1} \\ -\zeta L H_2 - \frac{r v_3 I H_2^2}{1+v_3 I H_2} \\ \frac{l v_1 H_1^2}{1+v_1 H_1} + \frac{l v_1 H_2^2}{1+v_1 H_2} + \frac{m v_2 L^2}{1+v_2 L} + \frac{m v_2 I^2}{1+v_2 I} \\ \frac{r v_3 L H_1^2}{1+v_3 L H_1} + \frac{r v_3 L H_2^2}{1+v_3 L H_2} + \frac{r v_3 I H_1^2}{1+v_3 I H_1} + \frac{r v_3 I H_2^2}{1+v_3 I H_2} \end{bmatrix}.$$

Here, $L_1 = H_1 + L_{TH_1} + I_{TH_1} + \eta_1(H_2 + L_{TH_2} + I_{TH_2}) + \eta_2(T_H + T_{TH})$ and $L_2 = I + I_{TH_1} + I_{TH_2}$. It can be easily noted that $\hat{H}(\mathcal{K}, \mathcal{L}) \not\leq 0$. Thus, the second condition for global stability of the equilibrium χ_0 is violated [33]. Therefore, perhaps the disease-free equilibrium χ_0 is not globally asymptotically stable.

6 Optimization problem

The main purpose of constructing and solving the optimization problem is to invent unrivalled strategy to restrain the HIV-TB dual epidemic, instead of applying baseline control technique. Our principal objective is to minimize the number of singly and dually infected individuals along with the cost of executing control measures in a specific time span $[0, t^*]$. Let $u_1(t)$, $u_2(t)$ and $u_3(t)$ be the control measures associated with the rates of consumption of HIV education, TB education and co-infection awareness, respectively. Further, let $u_4(t)$, $u_5(t)$ and $u_6(t)$ be the control functions relating to treatment rates, which are the fractions of additional infected people accessing treatments of HIV, TB and co-infection, respectively. The initial awareness consumption rates are considered as α_{E0}^H , α_{E0}^T and α_{E0}^{TH} ; initial treatment rates are assumed as l_0 , m_0 and r_0 . Furthermore, $\alpha_{E_{max}}^H$, $\alpha_{E_{max}}^T$ and $\alpha_{E_{max}}^{TH}$ indicate the

accordingly. After applying the aforementioned control measures, system (1) can be reformulated as follows:

$$\begin{aligned} \dot{S}_U^{TH} &= \Lambda - [(\alpha_E^H + u_1) + (\alpha_E^T + u_2) + (\alpha_E^{TH} + u_3) + \beta_T + \beta_H + \mu] S_U^{TH}, \\ \dot{S}_E^H &= (\alpha_E^H + u_1) S_U^{TH} - [\beta_1 \beta_H + \beta_T + (\alpha_E^T + u_2) + \mu] S_E^H, \\ \dot{S}_E^T &= (\alpha_E^T + u_2) S_U^{TH} - [\beta_2 \beta_T + \beta_H + (\alpha_E^H + u_1) + \mu] S_E^T + \rho T_T, \\ \dot{S}_E^{TH} &= (\alpha_E^{TH} + u_3) S_U^{TH} + (\alpha_E^T + u_2) S_E^H + (\alpha_E^H + u_1) S_E^T - (\beta_1 \beta_H + \beta_2 \beta_T + \mu) S_E^{TH}, \\ \dot{H}_1 &= \beta_H S_U^{TH} + \beta_1 \beta_H S_E^H + \beta_H S_E^T + \beta_1 \beta_H S_E^{TH} + \beta_H T_T - \phi \beta_T H_1 - (\delta + \mu) H_1 - \frac{(l + u_4) H_1}{1 + v_1 H_1}, \\ \dot{H}_2 &= \delta H_1 - (\phi \beta_T + \mu + d_H) H_2 - \frac{(l + u_4) H_2}{1 + v_1 H_2}, \\ \dot{L} &= \beta_T S_U^{TH} + \beta_T S_H^E + \beta_2 \beta_T S_E^T + \beta_2 \beta_T S_E^{TH} - (\theta \beta_H + \psi + \mu) L - \frac{(m + u_5) L}{1 + v_2 L}, \\ \dot{I} &= \psi L - (\theta \beta_H + \mu + d_T) I - \frac{(m + u_5) I}{1 + v_2 I}, \\ \dot{L}_{H_1} &= \theta \beta_H L + \phi \beta_T H_1 + \phi_1 \beta_T T_H \end{aligned}$$

$$\begin{aligned}
 & -(\sigma + \mu + \zeta)L_{H_1} \\
 & \frac{(r + u_6)L_{H_1}}{1 + v_3L_{H_1}}, \\
 \dot{L}_{H_2} = & \phi\beta_T H_2 + \sigma L_{H_1} \\
 & -(\zeta + d_H + \mu)L_{H_2} - \frac{(r + u_6)L_{H_2}}{1 + v_3L_{H_2}}, \\
 \dot{I}_{H_1} = & \theta\beta_H I + \zeta L_{H_1} - (\sigma + d_T + \mu)I_{H_1} \\
 & \frac{(r + u_6)I_{H_1}}{1 + v_3I_{H_1}}, \\
 \dot{I}_{H_2} = & \sigma I_{H_1} + \zeta L_{H_2} - (d_{TH} + \mu)I_{H_2} \\
 & \frac{(r + u_6)I_{H_2}}{1 + v_3I_{H_2}}, \\
 \dot{T}_T = & \frac{(m + u_5)L}{1 + v_2L} + \frac{(m + u_5)I}{1 + v_2I} \\
 & -\beta_H T_T - (\mu + \rho)T_T, \\
 \dot{T}_H = & \frac{(l + u_4)H_1}{1 + v_1H_1} \\
 & + \frac{(l + u_4)H_2}{1 + v_1H_2} - \phi_1\beta_T T_H - \mu T_H, \\
 \dot{T}_{TH} = & \frac{(r + u_6)L_{H_1}}{1 + v_3L_{H_1}} \\
 & + \frac{(r + u_6)L_{H_2}}{1 + v_3L_{H_2}} + \frac{(r + u_6)I_{H_1}}{1 + v_3I_{H_1}} \\
 & + \frac{(r + u_6)I_{H_2}}{1 + v_3I_{H_2}} - \mu T_{TH}. \tag{21}
 \end{aligned}$$

The optimization problem is proposed as

$$\begin{aligned}
 & \mathcal{J}(u_1^*, u_2^*, u_3^*, u_4^*, u_5^*, u_6^*) \\
 & = \min_{u \in \mathcal{U}} \mathcal{J}(u_1, u_2, u_3, u_4, u_5, u_6), \tag{22}
 \end{aligned}$$

where the objective function is given by,

$$\begin{aligned}
 \mathcal{J}(u) = & \int_0^{t^*} \left[A_1 H_1(t) + A_2 H_2(t) + A_3 I(t) \right. \\
 & + A_4 I_{H_1}(t) + A_5 I_{H_2}(t) \\
 & + B_1 u_4(t)(H_1(t) + H_2(t)) \\
 & + B_2 u_5(L(t) + I(t)) + B_3 u_6(L_{H_1}(t) \\
 & + L_{H_2}(t) + I_{H_1}(t) \\
 & + I_{H_2}(t)) + \frac{C_1}{2} u_1^2(t) + \frac{C_2}{2} u_2^2(t) \\
 & + \frac{C_3}{2} u_3^2(t) + \frac{C_4}{2} u_4^2(t) \\
 & \left. + \frac{C_5}{2} u_5^2(t) + \frac{C_6}{2} u_6^2(t) \right] dt \tag{23}
 \end{aligned}$$

regulated by the state system (21) with the set of permissible control measures \mathcal{U} such that

$$\mathcal{U} = \{(u_1(t), u_2(t), u_3(t), u_4(t), u_5(t), u_6(t))$$

$$\begin{aligned}
 & \in \mathbb{R}^6 | u_1(t), u_2(t), u_3(t), u_4(t), u_5(t), u_6(t) \\
 & \text{are Lebsgue integrable, } u_1(t) \in [0, \alpha_{E_{max}}^H - \alpha_{E_0}^H], \\
 & u_2(t) \in [0, \alpha_{E_{max}}^T - \alpha_{E_0}^T], \\
 & u_3(t) \in [0, \alpha_{E_{max}}^{TH} - \alpha_{E_0}^{TH}], \\
 & u_4(t) \in [0, l_{max} - l_0], \\
 & u_5(t) \in [0, m_{max} - m_0], \\
 & u_6(t) \in [0, r_{max} - r_0]\}.
 \end{aligned}$$

Here, $A_1 H_1$, $A_2 H_2$, $A_3 I$, $A_4 I_{H_1}$ and $A_5 I_{H_2}$ indicate the expenses for HIV-infected individuals without any symptom of AIDS, symptomatic people with AIDS, active TB-infected individuals, active TB-infected people who also have HIV, and AIDS-infected individuals who have co-infection of active TB, respectively. The cost of HIV treatment, TB treatment and HIV-TB co-infection treatment are represented by $B_1 u_4(t)(H_1(t) + H_2(t)) + \frac{C_4}{2} u_4^2(t)$, $B_2 u_5(L(t) + I(t)) + \frac{C_5}{2} u_5^2(t)$ and $B_3 u_6(L_{H_1}(t) + L_{H_2}(t) + I_{H_1}(t) + I_{H_2}(t)) + \frac{C_6}{2} u_6^2(t)$, respectively. $\frac{C_1}{2} u_1^2(t)$, $\frac{C_2}{2} u_2^2(t)$ and $\frac{C_3}{2} u_3^2(t)$ depict the outlay on HIV educational attempt, TB educational attempt and awareness attempt for co-infection, respectively.

Now, we employ Pontryagin’s Minimum Principal [34] to derive the necessary conditions for the existence of solution of optimization problem, that converts the optimization problem into minimization of the following Hamiltonian \mathcal{H} with respect to state trajectory x_i , optimal control u_i and corresponding Lagrange multiplier vector λ :

$$\begin{aligned}
 \mathcal{H} = & A_1 H_1(t) + A_2 H_2(t) + A_3 I(t) + A_4 I_{H_1}(t) \\
 & + A_5 I_{H_2}(t) + B_1 u_4(t)(H_1(t) + H_2(t)) \\
 & + B_2 u_5(L(t) + I(t)) + B_3 u_6(L_{H_1}(t) + L_{H_2}(t) \\
 & + I_{H_1}(t) + I_{H_2}(t)) + \frac{C_1}{2} u_1^2(t) + \frac{C_2}{2} u_2^2(t) \\
 & + \frac{C_3}{2} u_3^2(t) + \frac{C_4}{2} u_4^2(t) + \frac{C_5}{2} u_5^2(t) \\
 & + \frac{C_6}{2} u_6^2(t) + \lambda_1 \dot{S}_U^{TH} + \lambda_2 \dot{S}_E^H \\
 & + \lambda_3 \dot{S}_E^T + \lambda_4 \dot{S}_E^{TH} \\
 & + \lambda_5 \dot{H}_1 + \lambda_6 \dot{H}_2 + \lambda_7 \dot{L} \\
 & + \lambda_8 \dot{I} + \lambda_9 \dot{I}_{H_1} + \lambda_{10} \dot{I}_{H_2} \\
 & + \lambda_{11} \dot{I}_{H_1} + \lambda_{12} \dot{I}_{H_2} \\
 & + \lambda_{13} \dot{T}_T + \lambda_{14} \dot{T}_H + \lambda_{15} \dot{T}_{TH}, \tag{24}
 \end{aligned}$$

where λ_i is the i th component of Lagrange multiplier λ .

The first optimality condition is obtained as the minimization of \mathcal{H} with respect to optimal control measure u_i ($i = 1 - 6$). Therefore, we must get $\frac{\partial \mathcal{H}}{\partial u_i} = 0$. Thus, for the optimal control measures, we have

$$\begin{aligned} \hat{u}_1 &= \frac{1}{C_1} [(\lambda_1 - \lambda_2)S_U^{TH} + (\lambda_3 - \lambda_4)S_E^T], \\ \hat{u}_2 &= \frac{1}{C_2} [(\lambda_1 - \lambda_3)S_U^{TH} + (\lambda_2 - \lambda_4)S_E^H], \\ \hat{u}_3 &= \frac{1}{C_3} [(\lambda_1 - \lambda_4)S_U^{TH}], \\ \hat{u}_4 &= \frac{1}{C_4} \left[\frac{(\lambda_5 - \lambda_{14})H_1}{1 + \nu_1 H_1} + \frac{(\lambda_6 - \lambda_{14})H_2}{1 + \nu_1 H_2} - B_1(H_1 + H_2) \right], \\ \hat{u}_5 &= \frac{1}{C_5} \left[\frac{(\lambda_7 - \lambda_{13})L}{1 + \nu_2 L} + \frac{(\lambda_8 - \lambda_{13})I}{1 + \nu_2 I} - B_2(L + I) \right], \\ \hat{u}_6 &= \frac{1}{C_6} \left[\frac{(\lambda_9 - \lambda_{15})L_{H_1}}{1 + \nu_3 L_{H_1}} + \frac{(\lambda_{10} - \lambda_{15})L_{H_2}}{1 + \nu_3 L_{H_2}} + \frac{(\lambda_{11} - \lambda_{15})I_{H_1}}{1 + \nu_3 I_{H_1}} + \frac{(\lambda_{12} - \lambda_{15})I_{H_2}}{1 + \nu_3 I_{H_2}} - B_3(L_{H_1} + L_{H_2} + I_{H_1} + I_{H_2}) \right]. \end{aligned}$$

Hence, in the specified bounded interval, the optimal controls are given as

$$\begin{aligned} u_1^* &= \min\{\alpha_{E_{max}}^H, \max\{\alpha_{E_0}^H, \hat{u}_1\}\}, \\ u_2^* &= \min\{\alpha_{E_{max}}^T, \max\{\alpha_{E_0}^T, \hat{u}_2\}\}, \\ u_3^* &= \min\{\alpha_{E_{max}}^{TH}, \max\{\alpha_{E_0}^{TH}, \hat{u}_3\}\}, \\ u_4^* &= \min\{l_{max}, \max\{l_0, \hat{u}_4\}\}, \\ u_5^* &= \min\{m_{max}, \max\{m_0, \hat{u}_5\}\}, \\ u_6^* &= \min\{r_{max}, \max\{r_0, \hat{u}_6\}\}. \end{aligned}$$

The second condition of Pontryagin’s Minimum Principal gives the adjoint equations as $\frac{d\lambda_i}{dt} = -\frac{\partial \mathcal{H}}{\partial x_i}$, with the condition $\lambda_i(t^*) = 0$ ($i = 1 - 15$). Now, we can obtain the solution of the optimization problem by solving the above two mentioned conditions along with the state system (21).

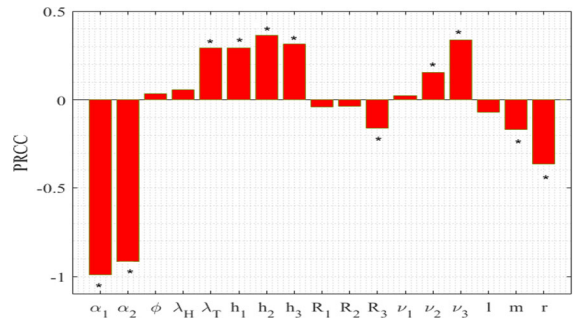


Fig. 2 Influence of uncertainty of the model (1) on individuals co-infected with active TB and HIV (I_{H_1}). Baseline values of parameters are the same as in Table 1. Model parameters having significant PRCCs are indicated by *

7 Numerical simulations

In this section, we perform extensive numerical simulations to explore the rich dynamics of model (1), and its submodels (2) and (10). We exemplify our analytical findings by commencing the initial conditions $S_U^{TH}(0) = 16500$, $S_E^H(0) = 80$, $S_E^T(0) = 60$, $S_E^{TH}(0) = 40$, $H_1(0) = 1250$, $H_2(0) = 250$, $L(0) = 9249$, $I(0) = 925$, $L_{H_1}(0) = 500$, $L_{H_2}(0) = 200$, $I_{H_1}(0) = 500$, $I_{H_2}(0) = 200$, $T_T(0) = 50$, $T_H(0) = 50$, $T_{TH}(0) = 58$ along with the parameter values given in Table 1. Treatment provision limitations necessitate the assumption of initial treatment rates as 0.62, 0.7 and 0.66 for HIV, TB and co-infection, respectively. Other parameter values are chosen within the epidemiologically feasible range and explained in previously published research papers [16,28,35]. Since the co-infection model (1) is a fifteen-dimensional system and hence it is recalcitrant while analyzing mathematically, thus we assess the full model numerically as much as possible.

7.1 Influence of some dominant parameters on HIV-TB co-infection

It is worthy to note that the parameter values taken for simulations may have some errors as they are obtained from several experiments. To tame this uncertainty, we perform global sensitivity analysis by implementing two statistical techniques: Latin Hypercube Sampling (LHS) and Partial Rank Correlation Coefficients (PRCCs) [36,37]. LHS permits to differ various parameters simultaneously in an efficient manner whereas the

latter one correlates the model parameters α_1 , α_2 , ϕ , λ_H , λ_T , h_1 , h_2 , h_3 , R_1 , R_2 , R_3 , ν_1 , ν_2 , ν_3 , l , m and r with the response function by assigning the values between -1 and 1 . We have taken the response function as co-infected class I_{H_1} . The signs of PRCCs indicate the type of correlation between the input parameters and the response function whereas the values represent the strength of their influences. We run 500 simulations per LHS by considering uniform distributions for each input parameter, and their baseline values are taken from Table 1 with $\pm 25\%$ deviations. We observe in Fig. 2 that the parameters ϕ , λ_H , λ_T , h_1 , h_2 , h_3 , ν_1 , ν_2 and ν_3 are positively correlated, and the parameters α_1 , α_2 , R_1 , R_2 , R_3 , l , m and r are negatively correlated with co-infected class I_{H_1} . Model parameters having significant PRCCs are marked specifically by *. The sensitivity results show that boosting up the propagation rates of information about HIV (α_1) and TB (α_2), and increasing the accessibility of treatment for co-infection by controlling the provision limitation have significant impacts in suppressing the burden of this dual epidemic.

In Figs. 3 and 4, we present sensitivity of awareness-induced basic reproduction numbers of HIV and TB submodels, respectively. In the figures, surface plots represent variations of these basic reproduction numbers with respect to two important model parameters at once. Figure 3 shows that the value of \mathcal{R}_E^H increases with the increase of HIV transmission rate (λ_H) and HIV to AIDS progression rate (δ). It can be diminished by reducing the probability of risky sexual practice (c_1) and time consumption parameter (h_1). The value of \mathcal{R}_E^H also decreases by increasing the HIV awareness distribution rate (α_1) and density of HIV awareness attempt (R_1). Similarly, Fig. 4 depicts that \mathcal{R}_E^T is positively associated with TB transmission rate (λ_T) and latent to active TB progression rate (ψ). Its value can be lessened by controlling the probability of risky sexual practice (c_2) and time consumption parameter (h_2). We also note that the value of \mathcal{R}_E^T diminishes with the increase of TB awareness distribution rate (α_2) and density of TB awareness attempt (R_2). These basic reproduction numbers can also be scaled down by increasing the treatment rates (l , m) and efficacy of awareness (ϵ). Thus, prevention control by proper educational attempt with less time consumption and treatment control with less limitation of provision can impose a restriction in disease transmission by suppressing the epidemic potential.

Next, to assess the influence of dominant parameters, we plot the singly infected individuals by differing two parameters simultaneously (see Fig. 5). In the figures, the contour lines signify the equilibrium values of singly infected populations. Figure 5a, b evince huge influence of time consumption parameter (h_1), provision limitation (ν_1) on HIV- and AIDS-infected population as HIV/AIDS-infected population increases with an increase in the values of h_1 and ν_1 . HIV infected populations (H_1 and H_2) are negatively associated with awareness dissemination rate (α_1) and treatment rate (l). Figure 5c, d shows the negative impacts of treatment rate (m) and TB awareness distribution rate (α_2) on latent TB-infected (L) and active TB-infected (I) individuals. It is apparent from the figure that on scaling up the time-consuming parameter (h_2) and resource-restricting parameter (ν_2), the number of TB-infected individuals increases massively. Overall, these figures depict the crucial roles of awareness and treatment in lowering the singly infected cases. It is necessary to augment the consumption of disease education by raising the awareness dissemination rates and minimizing the time consumption for educational attempts. Importantly, to curb the disease, medical support is the main weapon to restrict the co-infection and that should be accessible by reducing the limitation of resources as much as possible.

Now, we investigate the impacts of some ruling parameters specifically on dually infected populations, by plotting the model variables L_{H_1} , L_{H_2} , I_{H_1} and I_{H_2} with numerical variations of parameters in pairs viz. (ν_1, ν_3) , (h_3, ν_3) , (l, m) and (α_3, r) , Fig. 6. The sizes of co-infected populations are depicted by the surfaces at equilibrium, i.e., steady state or persevering oscillation. Presence of two surfaces in a figure indicates the highest and lowest numbers of co-infected individuals appearing in a limit cycle. In Fig. 6a, we assess the impact of provision limitation of HIV treatment as well as disruption in HIV-TB co-infection treatment on dual infection. It is clear from the figure that scaling up the resource limitation of HIV treatment induces disruption in co-infection treatment too, i.e., the value of ν_3 raises together with the value of ν_1 and minute limitation of HIV treatment produces a large number of co-infection, along with the increased restriction of co-infection treatment. The influence of time consumption in HIV-TB co-infection-related educational attempt and provision restriction of co-infection treatment are illustrated, reciprocally, in Fig. 6b. The fig-

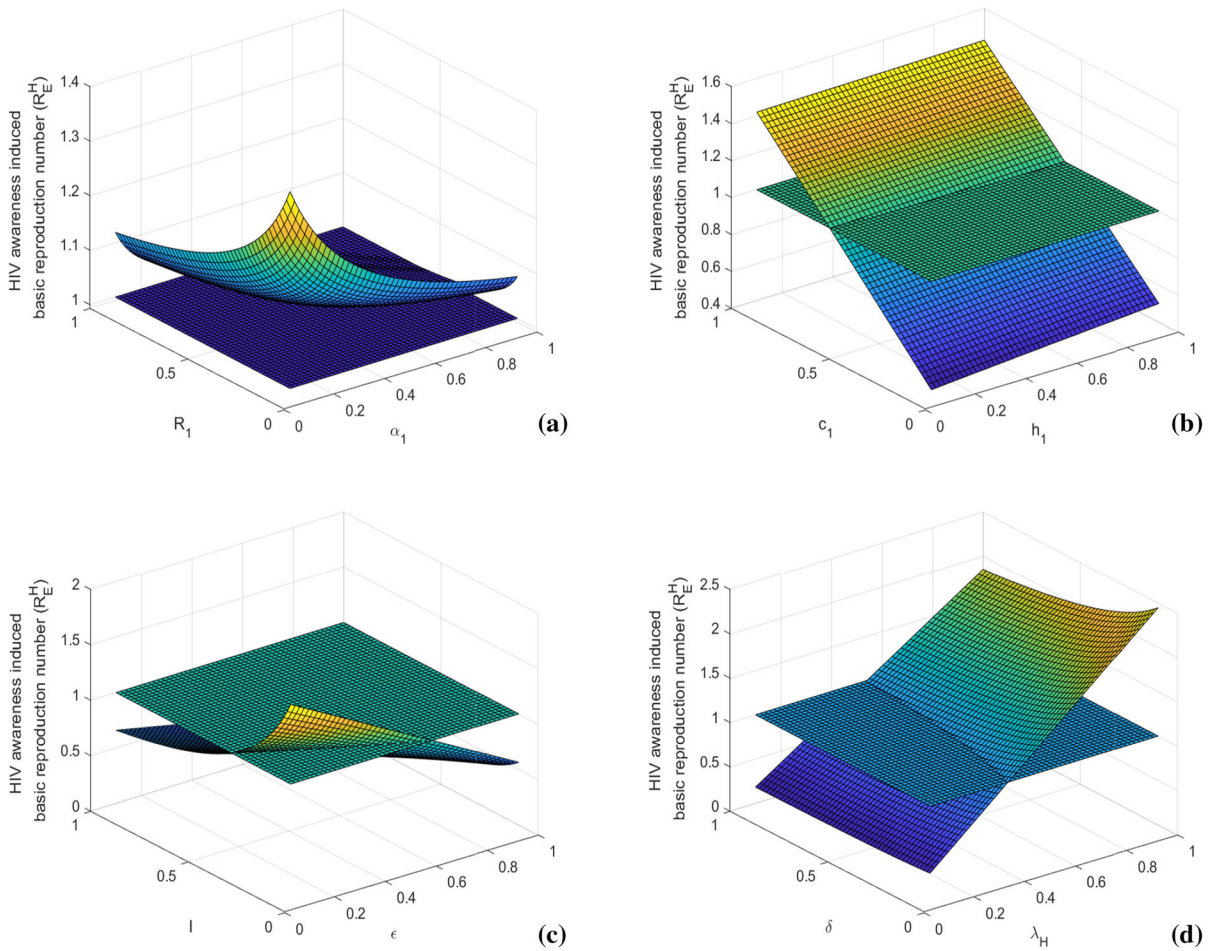


Fig. 3 Variations of awareness-induced basic reproduction number for HIV (\mathcal{R}_E^H) with respect to (a) α_1 and R_1 , (b) h_1 and c_1 , (c) ϵ and l , and (d) λ_H and δ . Rest of the parameters are at the same values as in Table 1

ures show that when the parameter h_3 surpasses the value 0.05, the resource limitation induces certainly an increase in the co-infected individuals. Co-infection also increases with the increase in values of (h_3, v_3) , which justifies the importance of awareness attempt in controlling dual-endemic, in a provision limitation condition. Figure 6c signifies that individual treatments of single infections impact the co-infection massively, by reducing the number of dually infected individuals. That is why, it is necessary to suppress the disruption in both HIV and TB medications. Figure 6d exhibits that co-infection diminishes rapidly with the growing treatment rate of co-infection (r) and HIV-TB awareness circulation rate (α_3).

7.2 Time series solutions and stability of equilibrium points

For parameter values in Table 1 and $v_1 = 0.00128$, we obtain the value of awareness-induced basic reproduction number for HIV model (2) as $\mathcal{R}_E^H = 0.6394$, which is less than unity. The HIV-free equilibrium is obtained as $\chi_0^H = (117988.85, 399578.77, 0, 0, 0)$, whose local stability is illustrated in Fig. 7a. The figure clearly shows the existence of locally asymptotically stable equilibrium of system (2). Further, Fig. 7b justifies Theorem 2 by illustrating the global stability of HIV-free equilibrium in $H_1 - H_2 - T_H$ space, when there is no provision restriction for HIV treatment, i.e., $v_1 = 0$. The solution trajectories plotted in Fig. 8a show the existence of a locally

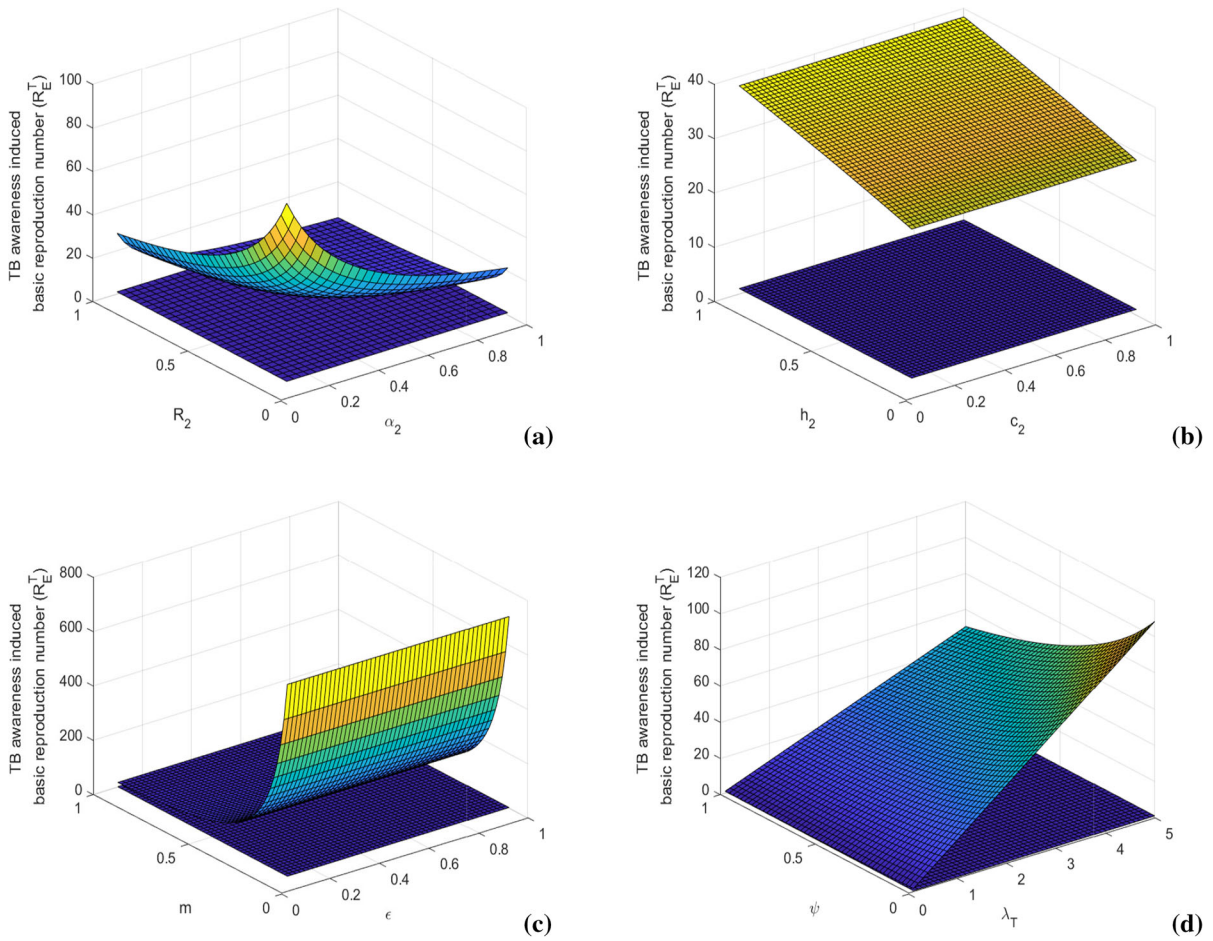


Fig. 4 Variations of awareness-induced basic reproduction number for TB (\mathcal{R}_E^T) with respect to (a) α_2 and R_2 , (b) h_2 and c_2 , (c) ϵ and m , and (d) λ_T and ψ . Rest of the parameters are at the same values as in Table 1

asymptotically stable HIV endemic equilibrium $\chi_H^1 = (31553.23, 12383.33, 74985.92, 21988.97, 2937.23)$ for $\nu_1 = 0.00129 > \bar{\nu}_1$, though $\mathcal{R}_E^H = 0.6394 < 1$. Figure 8b shows the existence of backward bifurcation by illustrating the co-existence of locally asymptotically stable endemic equilibrium χ_H^1 and stable DFE χ_0^H in $S_U^H - H_1 - T_H$ space. In Fig. 9a, we illustrate the existence of HIV endemic equilibrium $\chi_H^2 = (44723.23, 27230.68, 14881.77, 6021.62, 20903.14)$ for $\mathcal{R}_E^H = 1.696 > 1$ by choosing $\lambda_H = 0.75, l = 0.32$ and $\delta = 0.259$. The figure depicts local asymptotic stability of the equilibrium χ_H^2 when $\nu_1 = 0$ as the solution trajectories starting from different initial conditions converge to χ_H^2 . Importantly, we observe that for $\nu_1 < \bar{\nu}_1$, the endemic equilibrium χ_H^2 exists in an unstable mode whenever $\mathcal{R}_E^H < 1$ whereas this

equilibrium becomes stable whenever $\mathcal{R}_E^H > 1$. This demonstrates an exchange in the stability property of the equilibrium χ_H^2 and the presence of forward bifurcation at $\mathcal{R}_E^H = 1$ (see Fig. 9b). Thus, Figs. 8 and 9a, b illustrate Theorem 3 numerically. Figure 9c indicates the stability of HIV endemic equilibrium $\chi_H^3 = (14762.41, 2439.34, 35848.29, 29006.04, 33.31)$ for $\nu_1 = 0.06$.

In Fig. 10a, we plot the populations of TB-only dynamics for $\mathcal{R}_E^T = 0.3948 < 1$. The figure exhibits TB-free equilibrium $\chi_0^T = (134426.34, 99517.73, 0, 0, 0)$ for sufficiently small value of resource limitation parameter, i.e., $\nu_2 = 0.00023$. In Fig. 10b, we justify Theorem 5 numerically, by illustrating the global stability of TB-free equilibrium when there is no provision limitation in TB treatment (i.e., $\nu_2 = 0$). Fig-

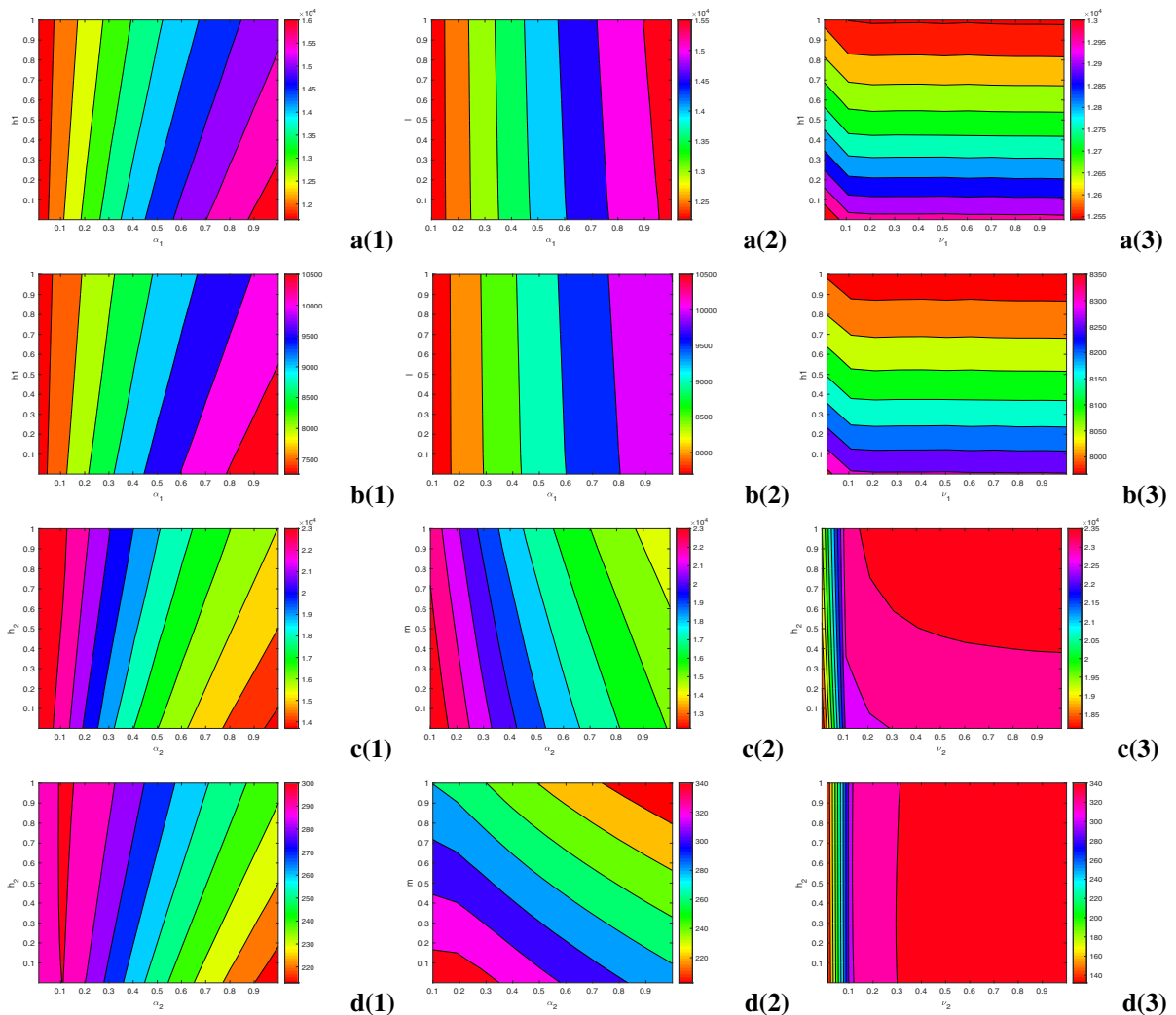


Fig. 5 Contour plots of HIV-infected individuals (first row) with respect to a(1) α_1 and h_1 , a(2) α_1 and l , a(3) v_1 and h_1 ; AIDS-infected individuals (second row) with respect to b(1) α_1 and h_1 , b(2) α_1 and l , b(3) v_1 and h_1 ; latent TB-infected individuals

(third row) with respect to c(1) α_2 and h_2 , c(2) α_2 and m , c(3) v_2 and h_2 ; and active TB-infected individuals (fourth row) with respect to d(1) α_2 and h_2 , d(2) α_2 and m , d(3) v_2 and h_2 . Other parameters are at the same values as in Table 1

ure 11a depicts that for $v_2 = 0.000355 > \bar{v}_2$, system (10) settles at the locally asymptotically stable endemic equilibrium $\chi_T^1 = (17300.17, 2729.57, 34221.08, 45293.17, 30652.07)$, in spite of having the basic reproduction number less than unity. Figure 11b shows the co-existence of locally asymptotically stable endemic equilibrium χ_T^1 and stable DFE χ_0^T for $\mathcal{R}_E^T < 1$ when $v_2 > \bar{v}_2$. For parameter values in Table 1 and $\lambda_T = 2.5, m = 0.35, \psi = 0.3$, we obtain $\mathcal{R}_E^T = 1.4933$. Figure 12a exhibits the existence of locally asymptotically stable TB endemic equilibrium $\chi_T^2 =$

(50241.19, 19454.27, 11717.93, 7477.29, 55984.71) for $v_2 = 0$. Figure 12a, b signifies the fact that the equilibrium χ_T^2 switches its stability at $\mathcal{R}_E^T = 1$ when $v_2 < \bar{v}_2$. Hence, Theorem 6 is illustrated numerically in Figs. 11 and 12a, b. Figure 12c reveals the stability of TB endemic equilibrium $\chi_T^3 = (5853.20, 340.96, 31910.63, 79729.32, 31.24)$ for $v_2 = 0.07$.

We plot the solution trajectories of the HIV-TB system (1) for different provision limitation conditions in Fig. 13. For the baseline parameter values in Table 1, we obtain $\mathcal{R}_E^H = 0.6394 < 1$ and $\mathcal{R}_E^T =$

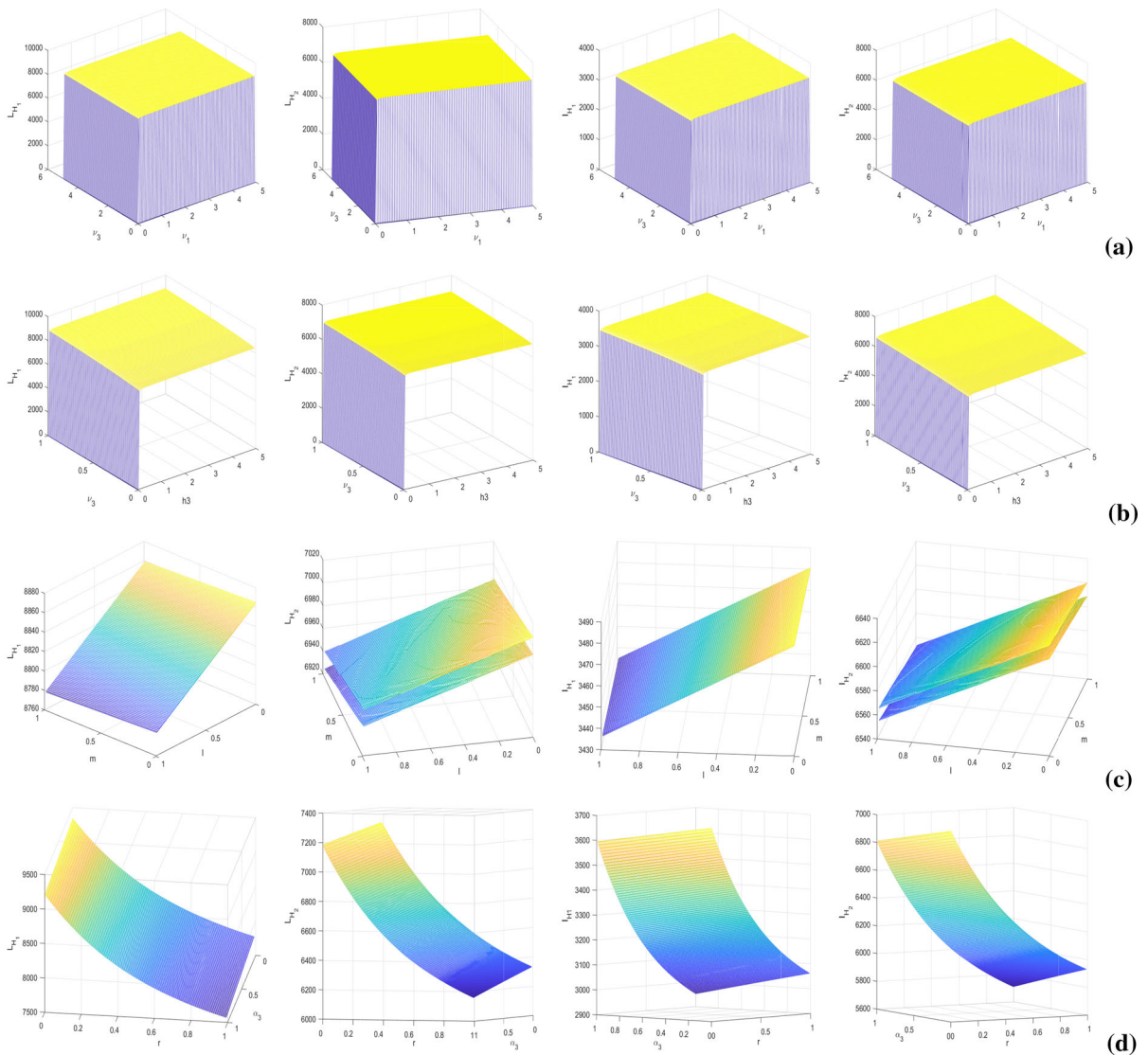


Fig. 6 Surface plots of co-infected individuals L_{H_1} (first column), L_{H_2} (second column), I_{H_1} (third column) and I_{H_2} (fourth column) with respect to (a) v_1 and v_3 , (b) h_3 and v_3 , (c) l and m , and (d) α_3 and r . Other parameters are at the same values as in Table 1

$0.3948 < 1$. Thus, $\mathcal{R}_E = 0.6394 < 1$. In Fig. 13(a), we can observe an stable HIV-TB endemic equilibrium $\chi_{HT}^1 = (9839.44, 804.82, 967.29, 2287.27, 2265.04, 213.11, 20458.72, 15427.09, 9409.04, 3717.76, 13077.21, 9973.91, 41.88, 20.32, 432.14)$ along with the disease-free equilibrium $\chi_0^1 = (42493.12, 37378.14, 31458.26, 423171, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0)$ under resource restriction condition, though the basic reproduction number is less than unity. In Fig. 13b, the solution trajectories of system (1) have been plotted in the absence of resource limitation for TB treat-

ment ($v_2 = 0$). In the figure, solid lines represent subpopulations in the presence of awareness attempt and the dashed lines indicate the individuals in the absence of TB awareness (i.e., $\alpha_E^T = 0, R_2 = 0, c_2 = 0$). It is apparent from the figure that the co-infection persists in the population when there is no educational attempt for TB, in spite of having no provision limitation of TB treatment. Though in this case, TB and co-infection are suppressed significantly up to a certain period of time, after that TB infection starts to increase with a rapid growth of co-infection

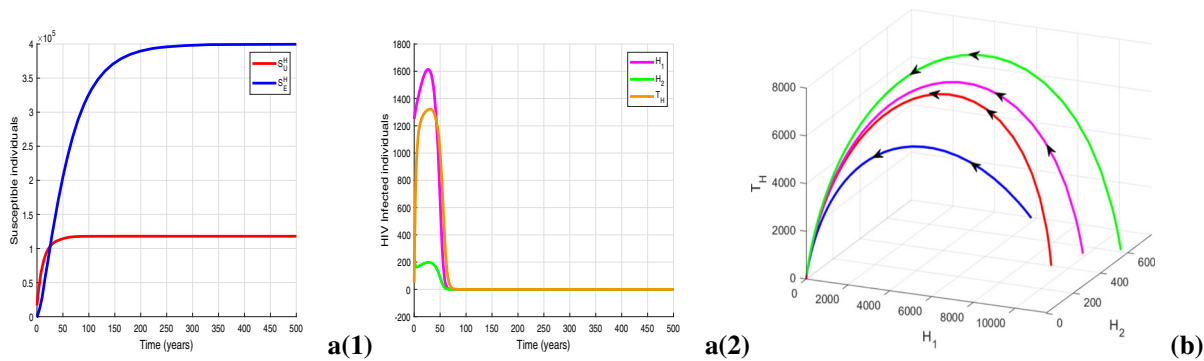


Fig. 7 Figures illustrating a(1)-a(2) HIV-free equilibrium χ_0^H with $\nu_1 = 0.00128$ for $\mathcal{R}_E^H = 0.6394$ and (b) globally asymptotically stable DFE with $\nu_1 = 0$. Other parameters are at the same values as in Table 1

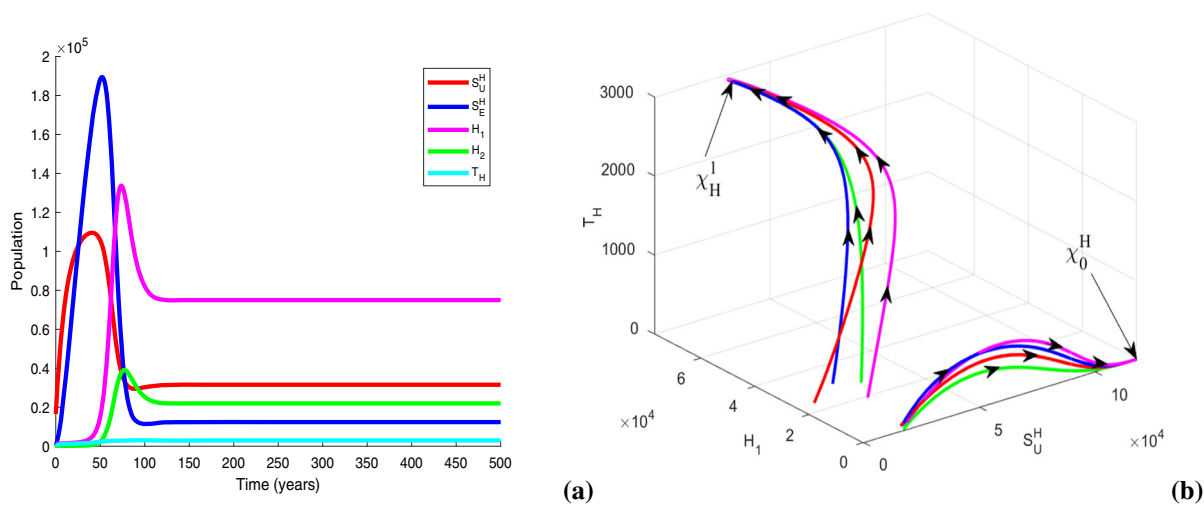


Fig. 8 (a) Solution trajectories of system (2) illustrating the existence of locally asymptotically stable HIV endemic equilibrium χ_H^1 with $\mathcal{R}_E^H = 0.6394 < 1$, $\nu_1 = 0.00129 > \bar{\nu}_1$ and (b) visual-

ization of backward bifurcation in $S_U^H - H_1 - T_H$ space. Other parameters are at the same values as in Table 1

and leads to an HIV-TB endemic equilibrium $\chi_{HT}^2 = (20229.09, 4371.03, 8058.40, 15879.69, 9367.06, 4648.26, 6694.92, 6694.92, 20593.44, 3921.5, 4225.19, 909.15, 7422.41, 40.37, 208.46)$. Moreover, on introducing TB awareness, TB infection and co-infection can be completely eradicated from the population, and there exists an HIV endemic equilibrium $\chi_H^4 = (21690.18, 6519.85, 3980.47, 16548.37, 70670.57, 22052.28, 0, 0, 0, 0, 0, 0, 0, 0, 0)$. This emphasizes the importance of awareness in resource limitation condition. In Fig. 13c, we plot the dynamics of system (1) for the situation when the HIV treatment provision limitation is zero ($\nu_1 = 0$). In this case, system (1) approaches to the HIV-TB endemic equilibrium $\chi_{HT}^3 =$

$(10022.68, 817.11, 998.37, 1983.69, 1170.01, 70.89, 21691.79, 17236.36, 8712.40, 3298.27, 12706.17, 9482.62, 43.14, 778.94, 178.84)$. Thus, we find that the HIV-TB co-infection persists in the population for inadequate treatment resources of both the diseases, in spite of having basic reproduction number below unity for HIV as well as TB submodels. Hence, these results reveal that it is not enough to scale down the basic reproduction number only, to curtail the co-infection. It is necessary to reduce the limitation for co-infection treatment by controlling HIV and TB both treatment restrictions to suppress the dual epidemic burden. However, the co-infection can be removed from the population by eliminating TB treatment provision limitation

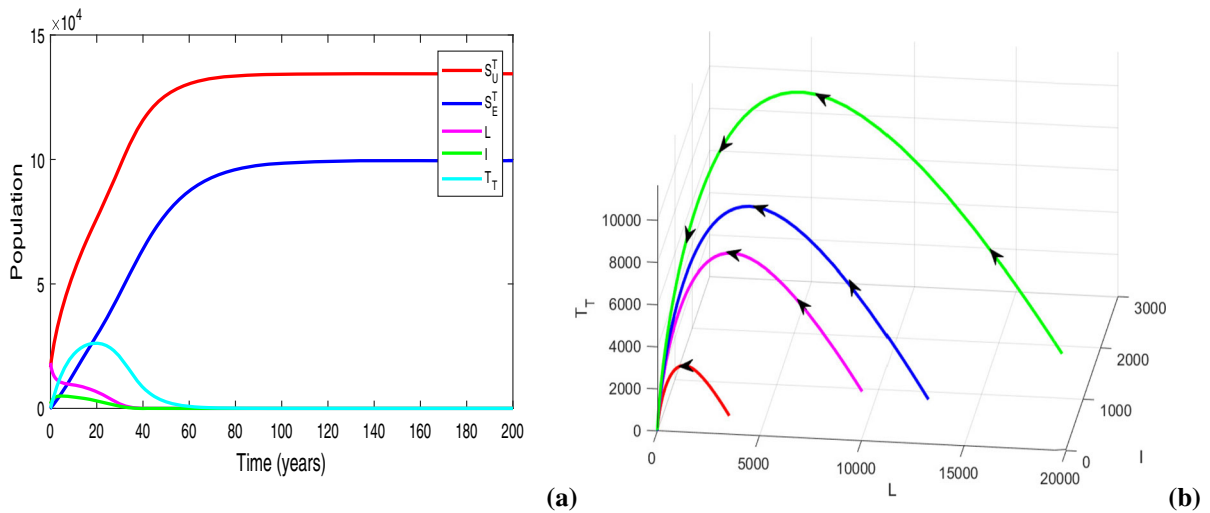


Fig. 10 Figures illustrating (a) TB disease-free equilibrium χ_0^T with $\nu_2 = 0.00023$ for $\mathcal{R}_E^T = 0.3948$, (b) globally asymptotically stable DFE with $\nu_2 = 0$. Other parameters are at the same values as in Table 1

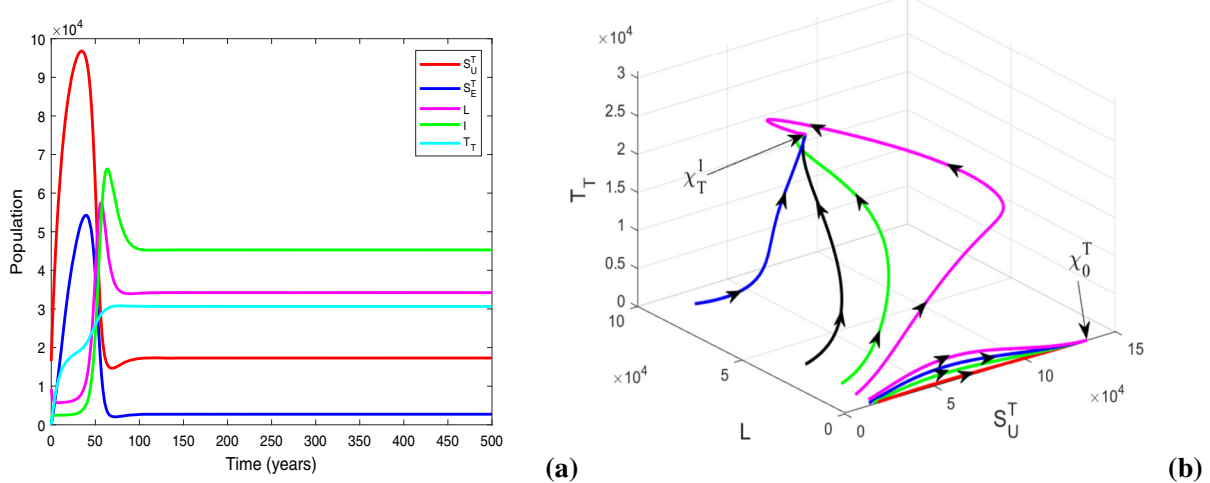


Fig. 11 (a) Solution trajectories of system (10) illustrating the existence of locally asymptotically stable TB endemic equilibrium χ_T^I with $\mathcal{R}_E^T = 0.3948 < 1$, $\nu_2 = 0.000355 > \bar{\nu}_2$, (b) visualization of backward bifurcation in $S_U^T - L - T_T$ space. Other parameters are at the same values as in Table 1

The figure generates the HIV-TB endemic equilibrium $\chi_{HT}^4 = (7848.65, 564.24, 494.52, 1360.08, 6163.59, 1836.55, 5922.91, 1798.37, 11887.24, 6810.22, 9522.91, 10834.97, 22.62, 12.72, 81.92)$. Further, Fig. 15b illustrates the global stability of HIV-TB endemic equilibrium χ_{HT}^4 .

7.3 Solution of optimization problem (21)

We implement fourth-order Runge–Kutta method to solve the proposed optimization problem (21) for a fixed time span of 10 years, by considering the afore-

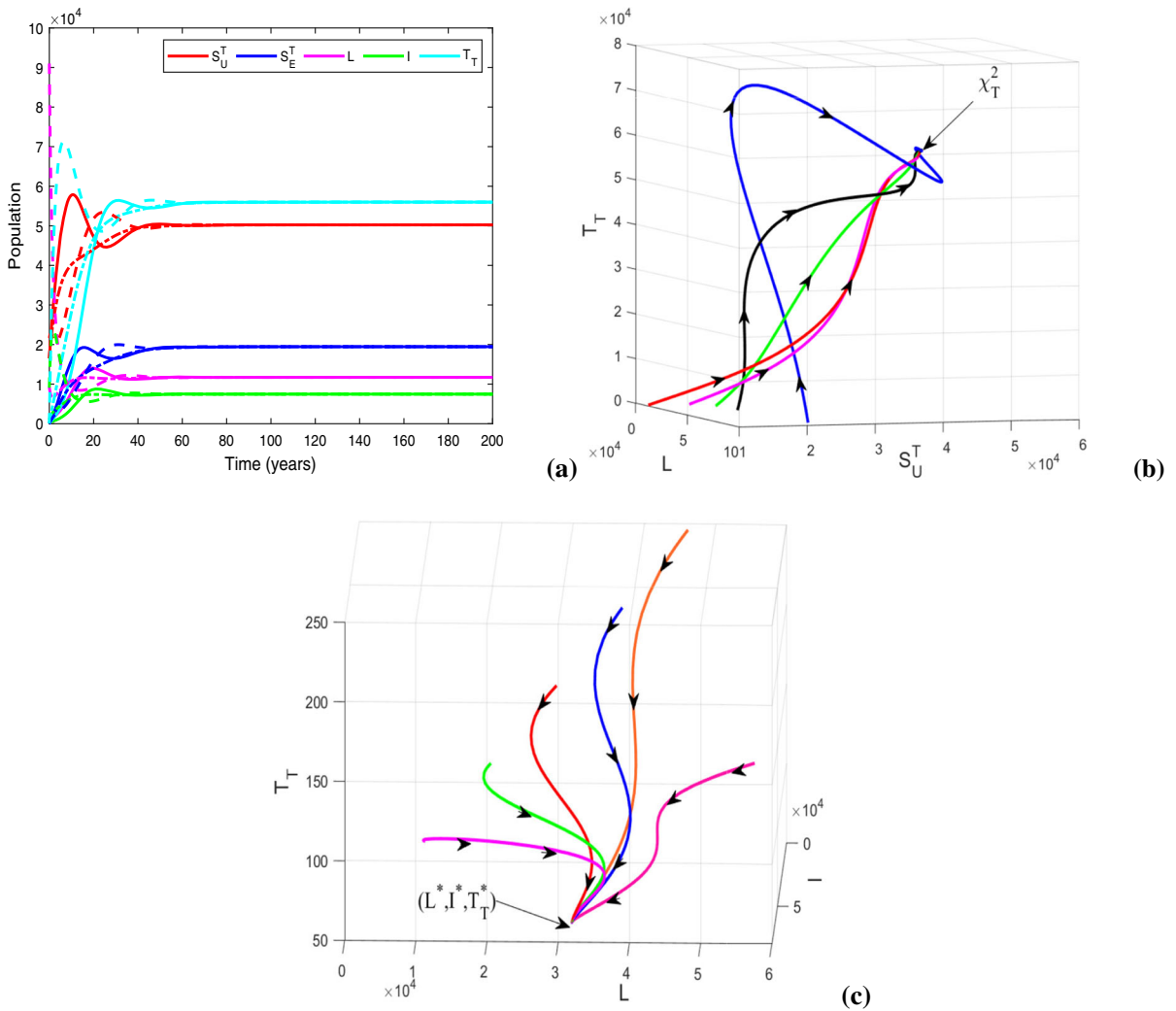


Fig. 12 (a) Solution trajectories of system (10) illustrating the existence of locally asymptotically stable TB endemic equilibrium χ_T^2 for $v_2 = 0$, (b) three-dimensional illustration of χ_T^2 and (c) global stability of another TB endemic equilibrium χ_T^3

in $L - I - T_T$ space for $v_2 = 0.07 > \bar{v}_2$ ($\mathcal{R}_E^H = 1.4933 > 1$ by taking $\lambda_T = 2.5$, $m = 0.35$ and $\psi = 0.3$). Other parameters are at the same values as in Table 1

mentioned initial conditions of the state variables. The optimization policy is taken into account for a specific period of time due to involvement of cost and limited provision in managing this dual-epidemic. The constants of the objective function are taken as $A_1 = 10$, $A_2 = 16$, $A_3 = 20$, $A_4 = 280$, $A_5 = 300$, $B_1 = 2$, $B_2 = 6$, $B_3 = 80$, $C_1 = 200$, $C_2 = 200$, $C_3 = 200$, $C_4 = 400$, $C_5 = 300$ and $C_6 = 800$. We initiate the process by guessing the controls satisfying the constraints. After that, the state equation system is solved beginning with the initial conditions onward and co-

state system is solved reversely. Figure 16 exhibits the influence of awareness control together with treatment control on the basic reproduction numbers \mathcal{R}_E^H and \mathcal{R}_E^T . In the figure, contour plots convey that the basic reproduction number can be diminished by applying these optimal techniques, that inspires us to construct such optimization problem.

Figure 17 demonstrates a comparison between baseline control case and optimal control case by illustrating the state trajectories. At the termination point of scheduled time span, there is a reduction in H_1 class

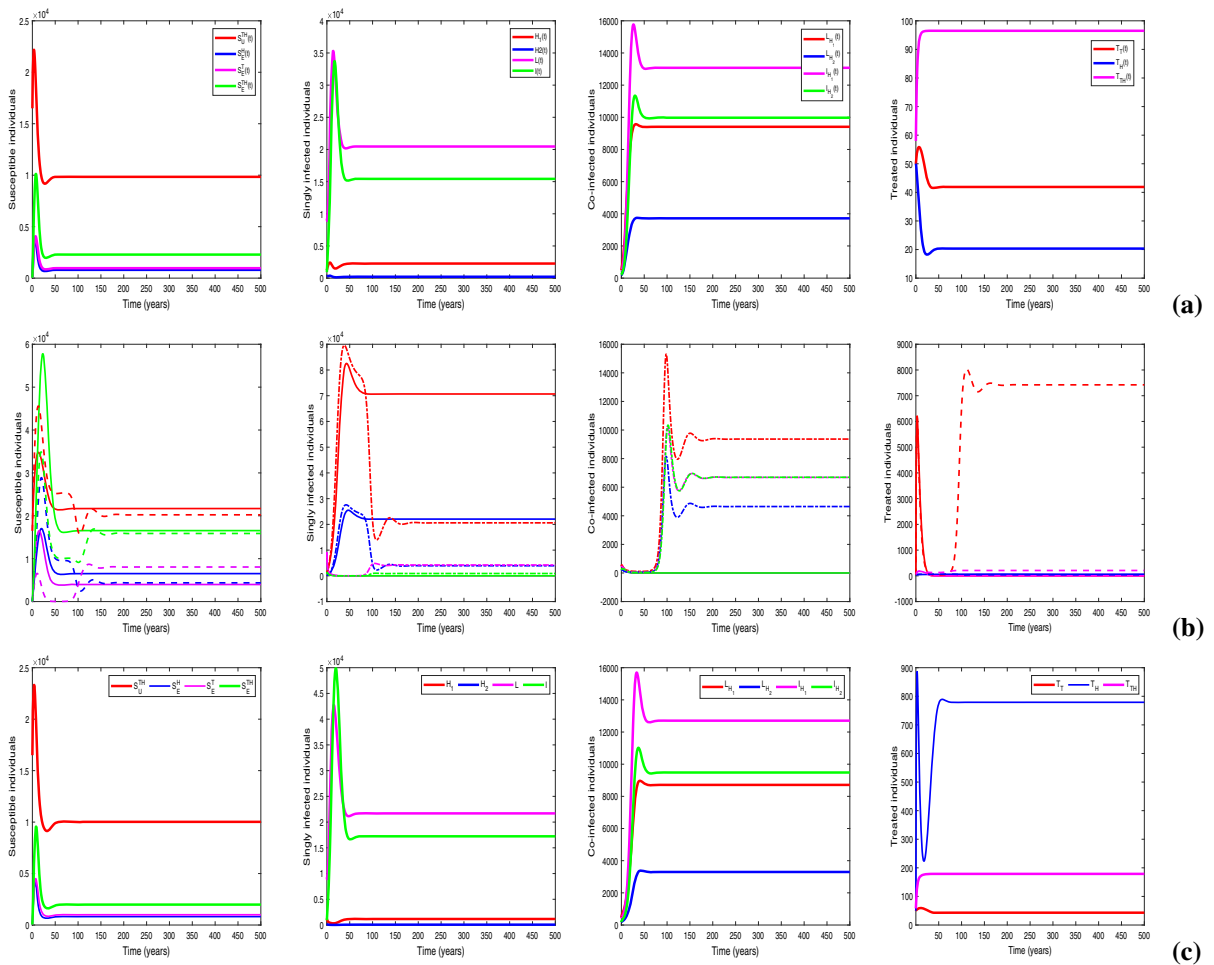


Fig. 13 Variations of susceptible, singly infected, co-infected and treated individuals with respect to time (a) under treatment provision limitation of HIV, TB and co-infection (i.e., $\nu_1 = 0.06$, $\nu_2 = 0.07$, $\nu_3 = 0.065$), (b) under treatment provision limitation of HIV and co-infection (i.e., $\nu_1 = 0.06$, $\nu_2 = 0$, $\nu_3 = 0.03$)

(The dashed lines represent the case of absence of TB awareness attempt, i.e., $\alpha_E^T = 0$, $R_2 = 0$, $c_2 = 0$), and (c) under treatment provision limitation of TB and co-infection (i.e., $\nu_1 = 0$, $\nu_2 = 0.07$, $\nu_3 = 0.035$). Other baseline parameter values are same as in Table 1

by 28.86%, in L class by 71.4%, in I class by 88.84%, in L_{H_1} class by 82.7%, in L_{H_2} class by 82.95%, in I_{H_1} class by 86.86% and in I_{H_2} class by 87.23%, in comparison with the case of baseline control technique. Thus, the proposed optimization mechanism has an enormous impact on both single and dual infections, that demolishes the epidemic curve under limited treatment provision condition induced by COVID-19 pandemic situation.

Figure 18 demonstrates this optimization approach, in which all the educational attempts and treatment controls are applied together for a planned time span of 10

years. As there is still no vaccination or curable medication for HIV and the HIV treatment structure is facing provision limitation, it is optimal to continue HIV awareness effort intensively for about 9.5 years from the starting. Initially, TB educational attempt should be applied in low intensity for almost 2.38 years, after that when TB infection becomes perceptible, it is necessary to execute TB awareness attempt with full intensity until 9.72 years. Co-infection awareness consumption is to be continued with a lower density throughout the planned time interval. The optimal strategy recommend to maintain 100% initial treatment rates for

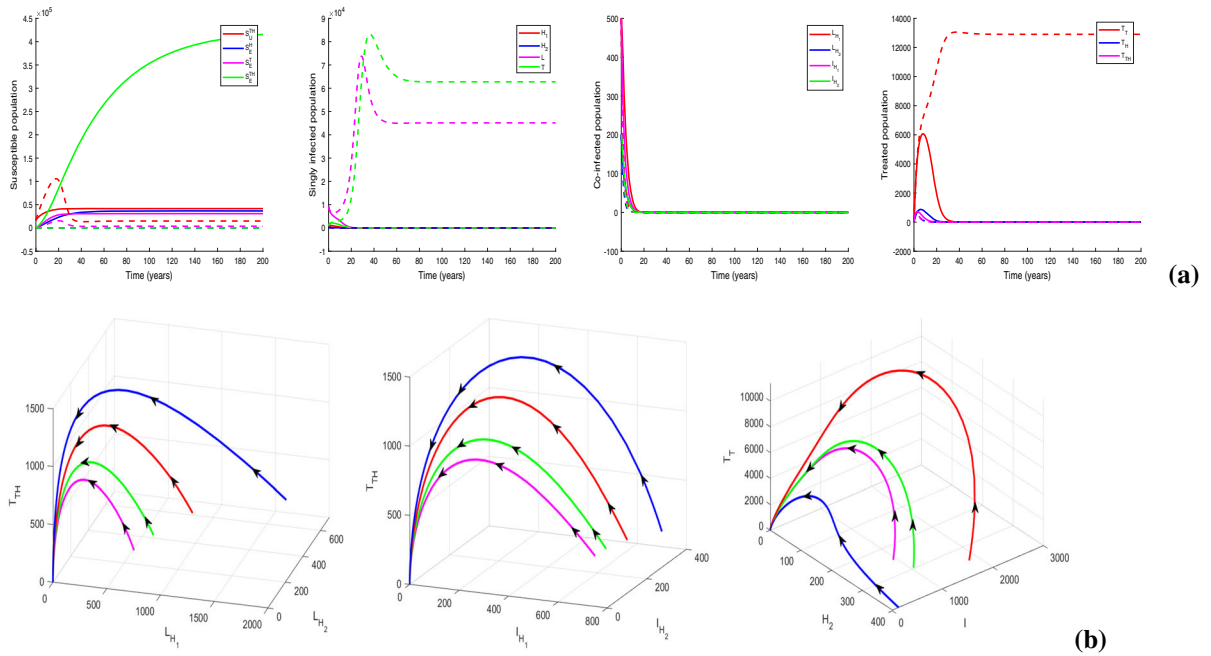


Fig. 14 Figures illustrating (a) HIV-TB disease-free equilibrium χ_0^2 (solid lines) and TB endemic equilibrium χ_T^4 in the absence of educational attempt (dashed lines) with $\nu_1 = 0.0011$, $\nu_2 = 0.00032$, $\nu_3 = 0.00142$ and (b) global stability of HIV-TB disease-free equilibrium when $\nu_1 = \nu_2 = \nu_3 = 0$. Other baseline parameter values are same as in Table 1

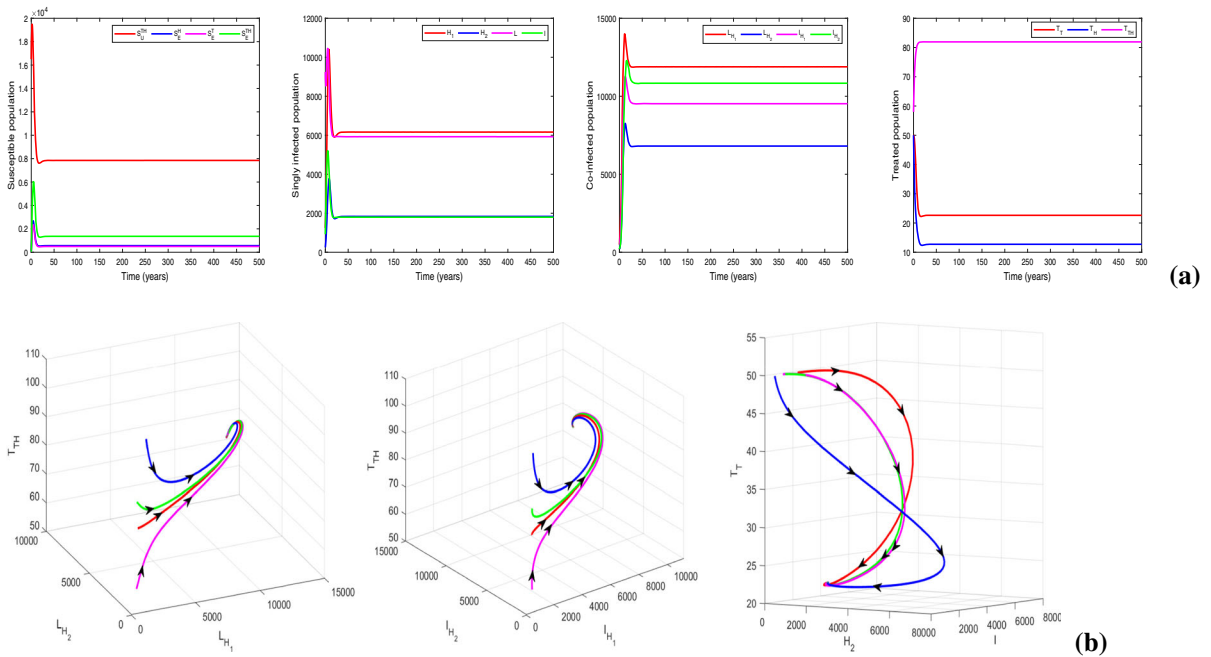


Fig. 15 Figures illustrating (a) HIV-TB endemic equilibrium χ_{HT}^4 for $\mathcal{R}_E = 1.69 > 1$, and (b) three-dimensional illustration of endemic equilibrium χ_{HT}^4 . Other baseline parameter values are same as in Table 1

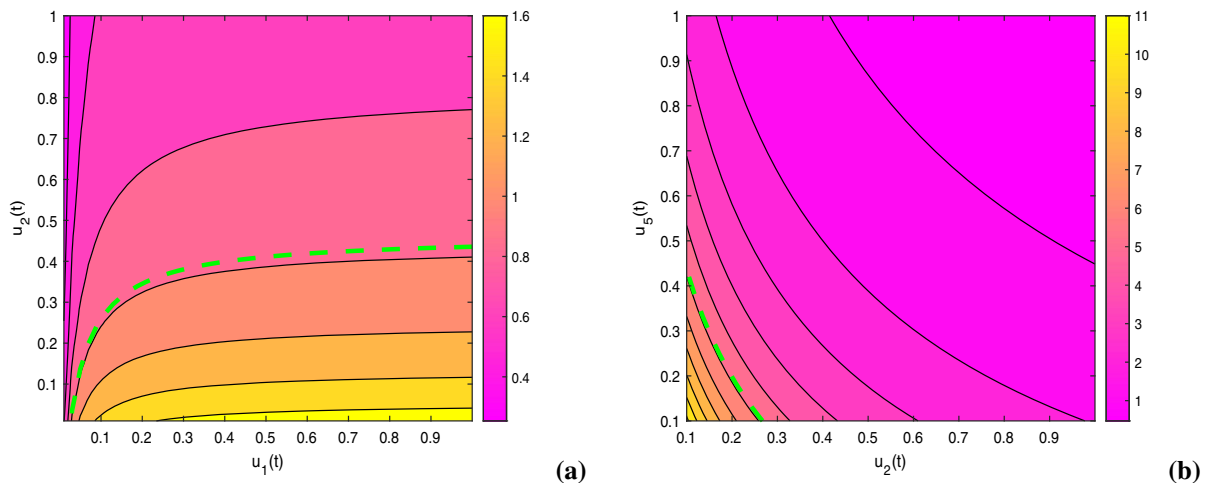


Fig. 16 Impact of optimal control measures on basic reproduction numbers (a) \mathcal{R}_E^H ($\alpha_1 = 0.0018$, $h_1 = 0.8$, $R_1 = 0.2$, $\lambda_H = 0.5$, $c_1 = 0.85$, $\epsilon = 0.0093$, $\delta = 0.59$, $l = 0.06$) and (b) \mathcal{R}_E^T . Other baseline parameter values are same as in Table 1

both HIV and TB individual infections until the end of planned time interval. However, the initial treatment effort for co-infected people should be continued for about 4 years. After that, it is ideal to reduce the initial treatment rate of co-infection till 8.83 years and uphold that decreased intensity up to the end of planned period.

Figure 19 depicts the variations in optimal control impact while curbing the dual-epidemic for different provision limitation conditions. In the figure, we have plotted the infected populations under optimization strategy for four cases viz. provision limitations for HIV, TB and co-infection ($v_1 = 0.06$, $v_2 = 0.07$, $v_3 = 0.065$); provision limitation for TB and co-infection ($v_1 = 0$, $v_2 = 0.07$, $v_3 = 0.035$); provision limitation for HIV and co-infection ($v_1 = 0.06$, $v_2 = 0$, $v_3 = 0.03$); and no provision limitation ($v_1 = v_2 = v_3 = 0$). We observe from the figure that the impact of optimal control is higher when there is no provision limitation for HIV and also in case of no resource limitation of TB treatment. Thus, it is deduced that as the resource restriction decreases, the impact of optimal control increases in restraining the outbreak. It can be inferred that optimal control has an immense effect in suppressing the dual pandemic burden if there is no provision restriction of HIV medication and has the greatest influence when there is no resource limitation for any treatment management.

8 Discussion

In this paper, an HIV-TB dual epidemic model that embraces provision limitation of treatments for both the diseases attributed to COVID-19 pandemic is investigated. In the proposed model, we have incorporated awareness attempts for both single as well as dual infections. The treatment rates are taken as Holling type-II functional pattern as the resources are limited for HIV, TB and co-infection medications. Also, we have taken into account four disjoint cohorts of susceptible individuals regarding the educational attempts of HIV, TB and co-infection, which makes our system more extensive. The model analysis reveals that awareness attempt significantly regulates the epidemic threshold. We found that for the model with either HIV or TB, the DFE is locally asymptotically stable whenever the awareness-induced basic reproduction number is below unity; further, it is globally asymptotically stable only when there is no provision limitation for treatment. We have also evaluated thresholds \bar{v}_1 and \bar{v}_2 for provision limitation parameters of the models with HIV and TB only, which are found to regulate the dynamical behavior of co-infection system. This assures the existence of locally asymptotically stable endemic equilibrium of each single infection model whenever the basic reproduction number is above unity provided the provision

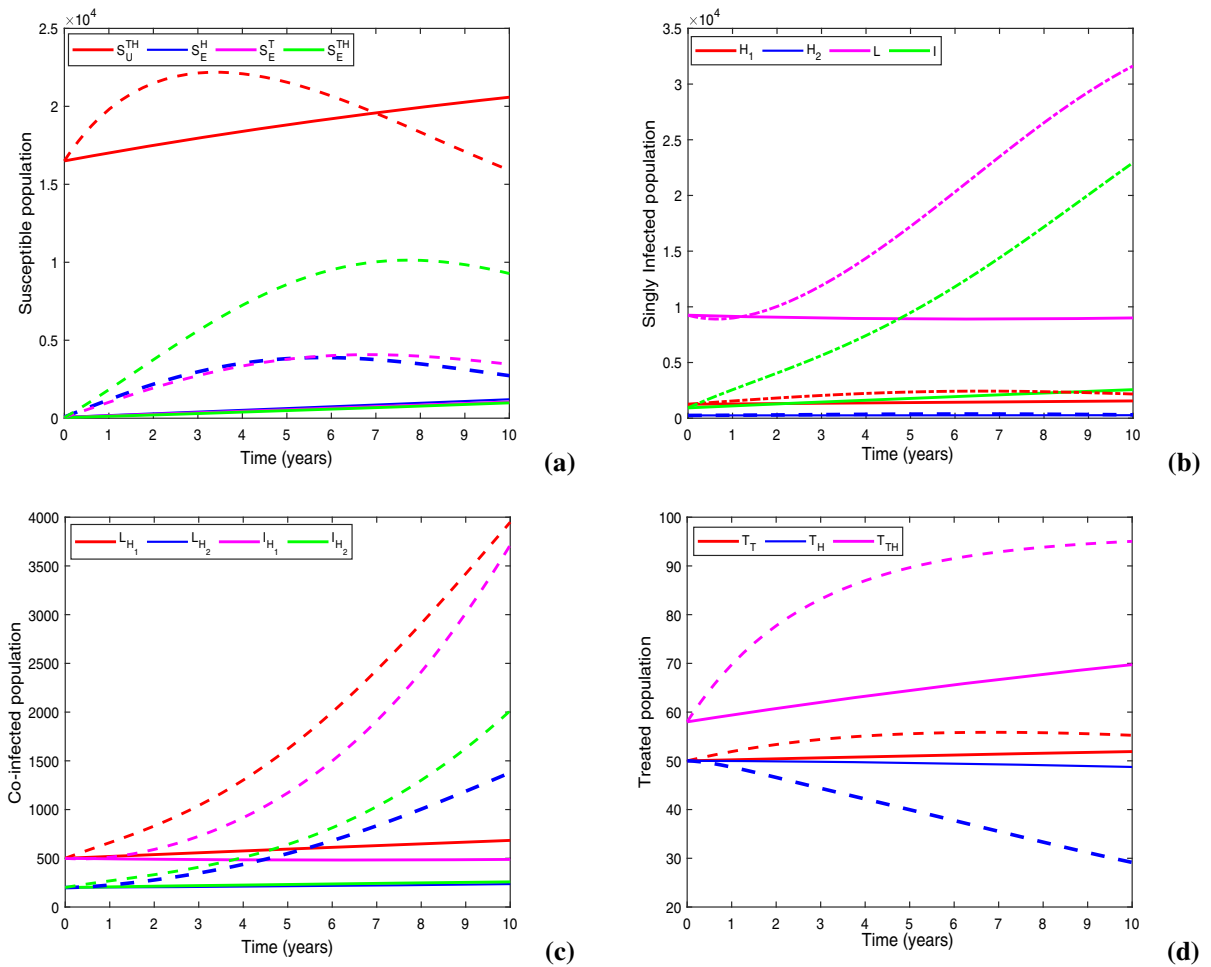


Fig. 17 Figures show the impacts of baseline control (dashed lines) and optimal control (solid lines) on (a) susceptible population, (b) singly infected population, (c) co-infected population and (d) treated population. Parameter values are same as in Table 1

limitation remains under the obtained threshold value. Moreover, the system undergoes a transcritical bifurcation while crossing the unit epidemic edge. If the basic reproduction number is below unity, the HIV and TB submodels are found to exhibit backward bifurcations provided the provision limitation parameters exceed their respective threshold values. For HIV-TB co-infection model, the DFE is locally asymptotically stable whenever $\mathcal{R}_E < 1$, but its global stability cannot be assured analytically. Overall, our analytical findings suggest to control the provision limitations for both the diseases to dominate the current situation.

Our numerical result explored the influences of dominant parameters on awareness-induced basic reproduction numbers. It recommends that epidemic aspect

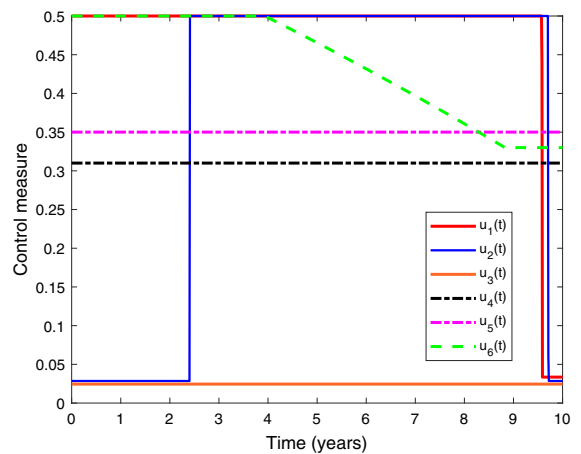


Fig. 18 Depiction of optimal control measures with respect to time. Parameter values are same as in Table 1

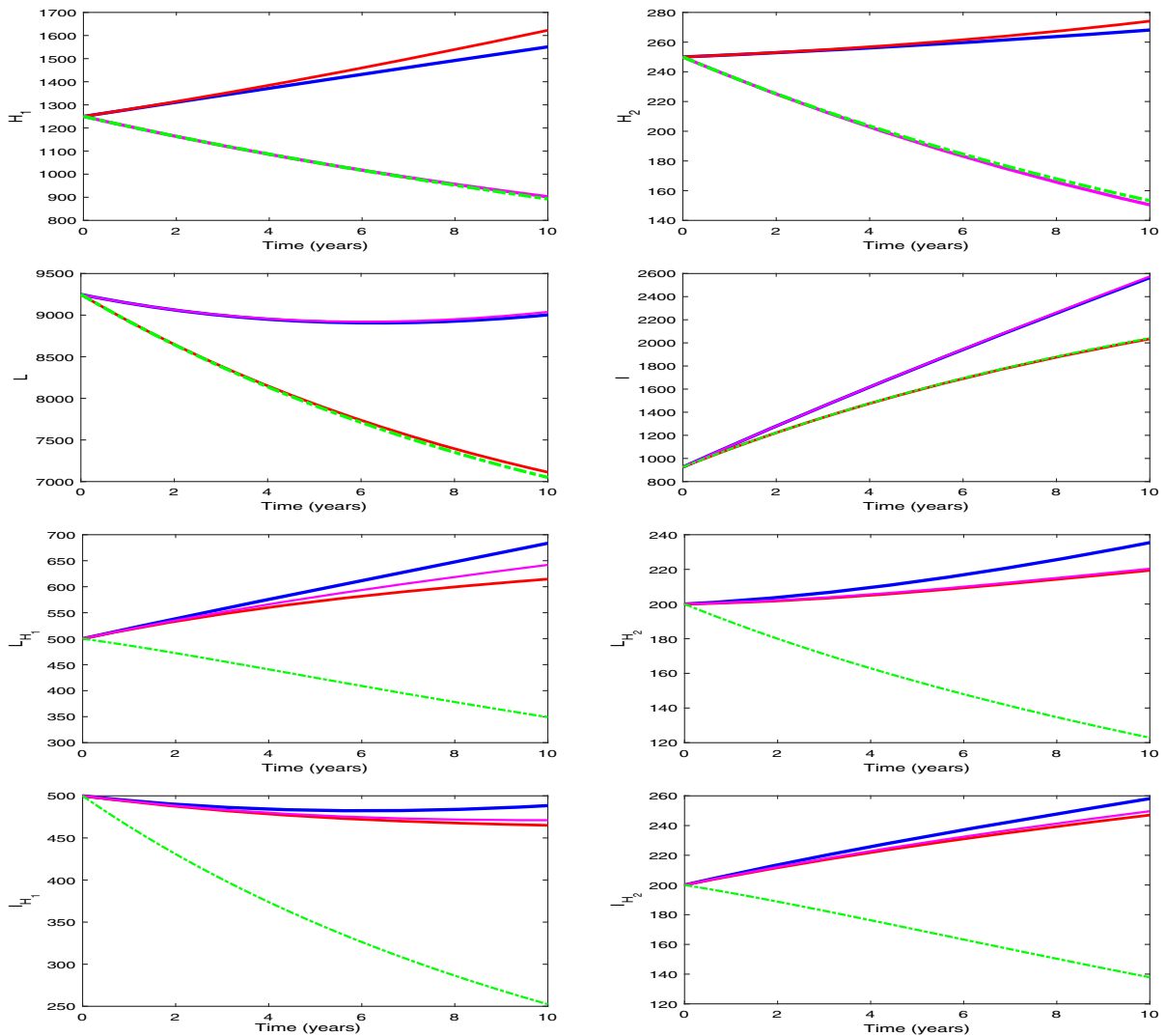


Fig. 19 Impacts of optimal control on infected individuals for $v_1 = 0.06, v_2 = 0.07, v_3 = 0.065$ (blue line); $v_1 = 0, v_2 = 0.07, v_3 = 0.035$ (magenta line); $v_1 = 0.06, v_2 = 0, v_3 = 0.03$ (red

line); and $v_1 = v_2 = v_3 = 0$ (green dash-dot line). Values of other parameters are the same as in Table 1

can be diminished by increasing the treatment rate along with consumption rate of awareness by scaling up the density of educational attempt, rate of propagating awareness and decreasing the consumption of time while making people aware. Assessment of the significant parameters' impact on infected individuals conveys the prominence of provision limitation that necessitates an urgent increase of treatment by curbing resource restriction to control the single as well as dual infections. We found that the endemic equilibrium exists under provision limitation situation for each sin-

gle disease model, though the basic reproduction number remains below unity. Limited treatment provisions for HIV, TB and co-infection exhibit the persistence of co-infection even though $\mathcal{R}_E < 1$. Therefore, it is not sufficient to scale down the basic reproduction number to curtail the dual epidemic; treatment accessibility should be increased by reducing the provision limitation. In case of basic reproduction number less than unity, in spite of eliminating the provision limitation of TB medication, co-infection relapses after a significant reduction and persists in the population for

a long time, if TB awareness is not propagated. But, it evinces HIV infection only under awareness control. Meanwhile, consideration of 100% accessibility of HIV treatment cannot eliminate the infection from the population. Therefore, co-infection can be curtailed by abolishing the provision limitation of TB treatment only if the awareness control is implemented. Moreover, when the provision restrictions are sufficiently small, TB infection remains in the population in the absence of any educational attempt in spite of having basic reproduction number less than unity. Although, analytical findings could not assure the global stability of DFE of the co-infection model, we have illustrated it numerically when there is no limitation of treatment provisions. We have also shown the global stability of endemic equilibrium of co-infection model numerically for $\mathcal{R}_E > 1$.

Finally, an optimization problem is constructed and solved that manifest epidemiological achievement by demolishing epidemic curve through optimization policy assimilating educational attempt along with treatment control of both single and dual infections, under limited medical provision. In the present provision limitation condition induced by COVID-19 pandemic, it is optimal to apply both awareness and treatment controls to curb the dual epidemic by following the strategy proposed in this study. Different impacts of our optimal policy for four individual resource restriction cases have been observed, exploring the vast influence of optimal control when resource of HIV treatment is not limited and also when TB service resource is not limited. It prescribes “no provision limitation” as the most impactful case while suppressing the co-infection load.

To the best of our knowledge, the model proposed in the present study is the first mathematical modeling approach to assess the current resource limitation condition of HIV and TB services induced by COVID-19 pandemic. Previously, a little bit attention has been paid on the impact of resource limitation on HIV-TB dual epidemic [28,35,38,39]. But, none of the previous study could consider the impact of COVID-19 situation. Sharomi et al. [38] investigated a model for HIV-TB co-infection by including four types of treatment strategies, suggesting that the HIV-only treatment policy is comparatively more effective than that of TB-only. They concluded that only one disease should be treated in case of resource limitation and universal strategy is the most beneficial for

co-infection control. Akwafuo et al. [39] studied the burden of HIV-TB co-infection in a low intervention area, West Africa, by presenting geographic analysis together with hybrid mathematical modeling to find the optimal control strategy. Their findings suggested that treatment of active TB-infected people before starting HIV treatment can reduce TB infection, and in resource limitation environment, testing and medication of TB disease should be accelerated for HIV negatives to prevent co-infection. Tanvi and Aggarwal [28] proposed an HIV-TB model introducing a constant time delay associated with retarded diagnosis and execution of treatment. They concluded that detection and treatment for both diseases within a proper time can result in economic and epidemic benefits. Authors did not consider saturated treatment rate in [28,38,39]. Later on, Tanvi et al. [35] introduced Holling type-II function in HIV-TB co-infection model as a resource-limited treatment rate of TB. But, they have not considered the matter of provision limitation in case of HIV treatment, and that is why, a constant treatment rate has been taken for HIV. Also, in [35], authors have not incorporated the awareness control. On the other hand, in the present investigation, we have incorporated Holling type-II treatment rates for both HIV and TB diseases by considering the provision limitation induced by COVID-19, and assessed the impact of awareness attempt under such condition. Our model assumptions exhibit more epidemiologically realistic and captivating dynamical behavior than the traditional epidemiological models and may contribute a lot to investigate the impact of recent provision restriction condition on HIV-TB dual epidemic. Moreover, the aforementioned studies could not provide any optimal policy for controlling the resource limitation situation. But, in the present study, we have introduced an optimization technique that gives a suitable strategy to control the dual epidemic by using awareness and treatment controls under the resource limitation condition. In [35], authors concluded that curbing TB infection can play a major role in controlling the co-infection under resource limitation condition for TB treatment. However, our findings provide specific thresholds of provision limitation that can help to diminish the restriction of HIV-TB medication. Further, our results suggest that increasing the accessibility of TB treatment by suppressing the limitation for TB service only is not enough to curb the dual epidemic under provision-restricted situation in absence of awareness. Our numerical findings convey that the co-

infection can persist in the population forever though the epidemic threshold is below unity if the current provision limitation situation continues in an unmanageable manner. Thus, awareness attempt should be introduced and provision limitation must be controlled for co-infection service to dominate the dual epidemic burden.

9 Conclusion

In this study, we have made an attempt for the first time to construct a mathematical model in order to investigate the present provision restriction condition of HIV and TB services induced by COVID-19 pandemic. Our findings suggest that, as the recent medical disruption caused by COVID-19 pandemic puts an enormous burden on the dual epidemic, the limitation of treatment provisions should be controlled as much as possible for HIV, TB and co-infection. Moreover, the crucial role of educational attempts is apparent from our investigation, in controlling the outbreak under provision restriction situation that include preventive mechanism, self-shielding measures, TB vaccination, motivation to acquire medication, etc. Most importantly, the derived thresholds for the treatment provision limitations of HIV and TB services can give a transparent idea about the extent to which the resource restriction should be diminished to curb the dual epidemic. Further, the optimization technique proposed in our study can be beneficial to construct practicable strategy to curtail HIV-TB dual epidemic with awareness and treatment control. This optimal policy will lead to epidemic as well as economic gain in this critical situation. The numerical results of the present investigation suggest that co-infection can be curtailed by suppressing provision restriction of TB treatment at the presence of awareness attempt and reducing the resource limitation for the treatment of co-infection is the best way to control the dual epidemic under resource restricted condition. Our results convey that the existing HIV-TB control programs need to be modified with recent situation-related awareness which will accelerate the patients' response to the healthcare system. According to the urgency of situation, people's behavior toward this dual epidemic must be modified as the responsible citizens. Necessary steps must be taken in order to make people aware about the consequence of COVID-19 impacting HIV-TB services. This should not be

neglected while fighting against COVID-19 pandemic, keeping the long-term effect in mind. The present mathematical approach may become the first step to catch the eyes on this matter.

The dynamics of COVID-19 has been assessed by numerous mathematical models since the starting of the pandemic [40,41]. In the present study, we did not incorporate the explicit dynamics of COVID-19 in the mathematical model formulation to investigate its direct influence on HIV-TB dual epidemic. Although our model explored an overall impact of resource limitation on HIV-TB dual epidemic due to COVID-19 pandemic situation, but to further extrapolate the results of this study, advance extension of our model system might be worth investigating. The present study can be extended by assimilating the dynamics of COVID-19. It would be interesting to examine the models with HIV and COVID-19, TB and COVID-19, and HIV, TB and COVID-19 co-infections individually.

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Data availability statements All data generated or analyzed during this study are included in this article.

Declarations

Conflict of interest The authors declare that there is no conflict of interests regarding the publication of this article.

Ethical standard The authors state that this research complies with ethical standards. This research does not involve either human participants or animals.

Appendix A

To investigate the global stability of disease-free equilibrium χ_0^H , we consider the following as a Lyapunov function candidate:

$$\mathcal{L}_1 = a_1 H_1 + a_2 H_2 + a_3 T_H,$$

where $a_1 = (\mu + l\eta_2)(l + \mu + d_H) + \delta(\mu\eta_1 + l\eta_2)$, $a_2 = (\mu\eta_1 + l\eta_2)(\delta + l + \mu)$, and $a_3 = \eta_2(\delta + l + \mu)(l + \mu + d_H)$. The Lyapunov derivative of \mathcal{L}_1 is obtained as,

$$\begin{aligned} \dot{\mathcal{L}}_1 &= a_1\dot{H}_1 + a_2\dot{H}_2 + a_3\dot{T}_H \\ &= a_1[\beta_H S_U^H + \beta_1\beta_H S_E^H - (\delta + \mu)H_1 - \tau_{H_1}] \\ &\quad + a_2[\delta H_1 - (\mu + d_H)H_2 - \tau_{H_2}] \\ &\quad + a_3(\tau_{H_1} + \tau_{H_2} - \mu T_H) \\ &= a_1 N\beta_H \left(\frac{\mu + \beta_1\alpha_E^H}{\mu + \alpha_E^H} \right) \\ &\quad - (H_1 + \eta_1 H_2 + \eta_2 T_H) \\ &\quad \left[\mu\left(\delta + \frac{l}{1 + \nu_1 H_1} + \mu\right)\left(\frac{l}{1 + \nu_1 H_2} + \mu + d_H\right) \right] \\ &= a_1 N\beta_H \left(\frac{\mu + \beta_1\alpha_E^H}{\mu + \alpha_E^H} \right) \\ &\quad - (H_1 + \eta_1 H_2 + \eta_2 T_H)[\mu(\delta + l + \mu)(l + \mu + d_H)], \text{ (if } \nu_1 = 0) \\ &= \left[\frac{N\beta_H\mu(\delta + l + \mu)(l + \mu + d_H)}{\lambda_H} \right] (\mathcal{R}_E^H - 1). \end{aligned}$$

Clearly, $\dot{\mathcal{L}}_1 \leq 0$ for $\mathcal{R}_E^H \leq 1$ only when the supply of treatment provisions has no limitation ($\nu_1 = 0$). As all the parameters of HIV model (2) are non-negative and $\mathcal{L}_1 = 0$ if and only if $H_1 = H_2 = T_H = 0$, so \mathcal{L}_1 is a Lyapunov function in Ω_1 and $\{\chi_0^H\}$ is the largest compact invariant set in $\{(S_U^H, S_E^H, H_1, H_2, T_H) \in \Omega_1 : \dot{\mathcal{L}}_1 = 0\}$. Consequently, by LaSalle’s Invariance Principle [31], when $\nu_1 = 0$, all the solutions of system (2) with initial conditions in Ω_1 proceed toward the HIV-free equilibrium χ_0^H as $t \rightarrow \infty$ for $\mathcal{R}_E^H \leq 1$. Therefore, if there is no limitation of treatment provisions, HIV infection will be eliminated from the population.

Appendix B

We apply the center manifold theory [32] to investigate the stability of endemic equilibrium point χ_H^* as standard linearization of system (2) around this equilibrium is strenuous and not persuadable mathematically. To use Theorem 4.1 of [32], we reconstruct the HIV model (2) in the form $\frac{dX}{dt} = f = [f_1, f_2, f_3, f_4, f_5]^T$, by taking $S_U^H = x_1, S_E^H = x_2, H_1 = x_3, H_2 = x_4, T_H = x_5$, and by introducing $X = [x_1, x_2, x_3, x_4, x_5]^T$ as follows:

$$\begin{aligned} \frac{dx_1}{dt} &= f_1 = \Lambda - (\alpha_E^H + \beta_H + \mu)x_1, \\ \frac{dx_2}{dt} &= f_2 = \alpha_E^H x_1 - (\beta_1\beta_H + \mu)x_2, \\ \frac{dx_3}{dt} &= f_3 = \beta_H x_1 + \beta_1\beta_H x_2 - (\delta + \mu)H_1 - \tau_{x_3}, \\ \frac{dx_4}{dt} &= f_4 = \delta x_3 - (\mu + d_H)x_4 - \tau_{x_4}, \\ \frac{dx_5}{dt} &= f_5 = \tau_{x_3} + \tau_{x_4} - \mu x_5. \end{aligned} \tag{25}$$

For $\mathcal{R}_0^H = 1$, we get a critical value of λ_H (say λ_H^*) as

$$\lambda_H^* = \frac{\mu(\mu + \alpha_E^H)(\delta + l + \mu)(l + \mu + d_H)}{(\mu + \beta_1\alpha_E^H)[\mu(l + \mu + d_H) + \delta(\mu\eta_1 + l\eta_2) + l\eta_2(l + \mu + d_H)]}. \tag{26}$$

Evaluating the Jacobian matrix of system (25) at DFE χ_H^0 , we get the following matrix

$$J_{\chi_0^H} = \begin{bmatrix} -(\alpha_E^H + \mu) & 0 & \frac{\mu\lambda_H^*}{\alpha_E^H + \mu} & \frac{\eta_1\mu\lambda_H^*}{\alpha_E^H + \mu} & \frac{\eta_2\mu\lambda_H^*}{\alpha_E^H + \mu} \\ \alpha_E^H & -\mu & \frac{\beta_1\lambda_H^*\alpha_E^H}{\alpha_E^H + \mu} & \frac{\beta_1\eta_1\lambda_H^*\alpha_E^H}{\alpha_E^H + \mu} & \frac{\beta_1\eta_2\lambda_H^*\alpha_E^H}{\alpha_E^H + \mu} \\ 0 & 0 & \left[\lambda_H^* \left(\frac{1 + \beta_1\alpha_E^H}{\alpha_E^H + \mu} \right) - (\delta + l + \mu) \right] & \left[\eta_1\lambda_H^* \left(\frac{1 + \beta_1\alpha_E^H}{\alpha_E^H + \mu} \right) \right] & \left[\eta_2\lambda_H^* \left(\frac{1 + \beta_1\alpha_E^H}{\alpha_E^H + \mu} \right) \right] \\ 0 & 0 & \delta & -(\mu + l + d_H) & 0 \\ 0 & 0 & a & a & -\mu \end{bmatrix}.$$

As $\lambda_H = \lambda_H^*$, the linearization matrix has a simple zero eigenvalue. Hence, the center manifold theory can be used to analyze the local asymptotic stability of the endemic equilibrium point χ_H^* .

There must exist a right eigenvector $w = [w_1, w_2, w_3, w_4, w_5]^T$ and a left eigenvector $v = [v_1, v_2, v_3, v_4, v_5]$ of Jacobian matrix $J_{\chi_0^H}$, associated with zero eigenvalue whose components at $\lambda_H = \lambda_H^*$ are obtained as

$$\begin{aligned} w_1 &= -\frac{\mu\lambda_H^*(w_3 + \eta_1w_4 + \eta_2w_5)}{(\alpha_E^H + \mu)^2}, \\ w_2 &= \frac{\beta_1\alpha_E^H\lambda_H^*(w_3 + \eta_1w_4 + \eta_2w_5)}{\mu(\alpha_E^H + \mu)^2} \\ +\alpha_E^H w_1, w_3 &> 0, w_4 = \frac{\delta w_3}{\mu + d_H + l}, \\ w_5 &= \frac{aw_3}{\mu} \left(1 + \frac{\delta}{\mu + d_H + l} \right); \\ \text{and } v_1 = v_2 = 0, v_3 &> 0, \\ v_4 &= \left(\frac{1}{\mu + l + d_H} \right) \\ \left[\frac{v_3\eta_1\lambda_H^*(\mu + \beta_1\alpha_E^H)}{\alpha_E^H + \mu} + lv_5 \right], \\ v_5 &= \frac{v_3\eta_2\lambda_H^*(\mu + \beta_1\alpha_E^H)}{\alpha_E^H + \mu}. \end{aligned}$$

Now, we compute the following second-order non-vanishing partial derivatives of f at the disease-free equilibrium point χ_H^0 to calculate a and b , following Theorem 4.1 of [32] (as $v_1 = v_2 = 0$, so there is no need to calculate the partial derivatives of f_1 and f_2), and get

$$\begin{aligned} \frac{\partial^2 f_3}{\partial x_3^2} &= 2av_1 - \frac{2\lambda_H^*\mu(\mu + \beta_1\alpha_E^H)}{\Lambda(\alpha_E^H + \mu)}, \\ \frac{\partial^2 f_3}{\partial x_3\partial x_4} &= -\frac{\mu\lambda_H^*(1 + \eta_1)(\mu + \beta_1\alpha_E^H)}{\Lambda(\alpha_E^H + \mu)}, \\ \frac{\partial^2 f_3}{\partial x_3\partial x_5} &= -\frac{\mu\lambda_H^*(1 + \eta_2)(\mu + \beta_1\alpha_E^H)}{\Lambda(\alpha_E^H + \mu)}, \end{aligned}$$

$$\begin{aligned} \frac{\partial^2 f_3}{\partial x_4^2} &= -\frac{2\mu\lambda_H^*\eta_1(\mu + \beta_1\alpha_E^H)}{\Lambda(\alpha_E^H + \mu)}, \\ \frac{\partial^2 f_3}{\partial x_4\partial x_5} &= -\frac{\mu\lambda_H^*(\eta_1 + \eta_2)(\mu + \beta_1\alpha_E^H)}{\Lambda(\alpha_E^H + \mu)}, \\ \frac{\partial^2 f_3}{\partial x_5^2} &= -\frac{2\mu\lambda_H^*\eta_2(\mu + \beta_1\alpha_E^H)}{\Lambda(\alpha_E^H + \mu)}, \\ \frac{\partial^2 f_4}{\partial x_5^2} &= 2lv_1, \frac{\partial^2 f_3}{\partial x_3\partial \lambda_H^*} = \frac{(\mu + \beta_1\alpha_E^H)}{\alpha_E^H + \mu}, \\ \frac{\partial^2 f_3}{\partial x_4\partial \lambda_H^*} &= \eta_1 \frac{(\mu + \beta_1\alpha_E^H)}{\alpha_E^H + \mu}, \\ \frac{\partial^2 f_3}{\partial x_5\partial \lambda_H^*} &= \eta_2 \frac{(\mu + \beta_1\alpha_E^H)}{\alpha_E^H + \mu}. \end{aligned}$$

By substituting the above values in the expressions for a and b , we have

$$\begin{aligned} a &= \sum_{k,i,j=1}^n v_k w_i w_j \frac{\partial^2 f_k}{\partial x_i \partial x_j}(0, 0) \\ &= 2lv_1 \left[v_3 w_3^2 + \left(\frac{v_4 w_4 \delta}{\mu + d_H + l} \right) \right] \\ &\quad - \frac{\lambda_H^* \mu v_3 w_3^2 (\mu + \beta_1 \alpha_E^H)}{\Lambda(\alpha_E^H + \mu)} \\ &\quad \left[2 + \frac{\delta(\eta_1 + 1)}{(\mu + l + d_H)} \right. \\ &\quad + \frac{\delta a(\eta_1 + \eta_2)}{\mu(\mu + l + d_H)} \left(1 + \frac{\delta}{\mu + l + d_H} \right) \\ &\quad + \frac{2a^2}{\mu^2} \left(1 + \frac{\delta}{\mu + l + d_H} \right)^2 \\ &\quad + 2\eta_1 \left(\frac{\delta}{\mu + l + d_H} \right)^2 \\ &\quad \left. + \frac{\delta l(1 + \eta_2)}{\mu(\mu + l + d_H)} \left(1 + \frac{\delta}{\mu + l + d_H} \right) \right], \\ b &= \sum_{k,i=1}^n v_k w_i \frac{\partial^2 f_k}{\partial x_i \partial \lambda_H^*}(0, 0) \end{aligned}$$

$$= \frac{v_3(\mu + \beta_1 \alpha_E^H)}{\alpha_E^H + \mu} (w_3 + \eta_1 w_4 + \eta_2 w_5) > 0.$$

It is clear that b is always positive. Note that $a < 0$ if $0 \leq v_1 < \bar{v}_1$, and $a > 0$ if the inequality is reversed. Therefore, by Theorem 4.1 of [32], it can be concluded that the endemic equilibrium point χ_H^* of HIV-only model (2) is locally asymptotically stable for $\mathcal{R}_0^H > 1$ with \mathcal{R}_0^H near 1 (i.e., $\lambda_H^* \approx \lambda_H$) and $0 \leq v_1 < \bar{v}_1$. Here, \bar{v}_1 is obtained as

$$\begin{aligned} \bar{v}_1 = & \frac{v_3 \mu (\mu + \beta_1 \alpha_E^H) (\mu + l + d_H)}{2l \Lambda (\alpha_E^H + \mu) [v_3 (\mu + l + d_H) + v_4 \delta]} \\ & \left[2 + \left(\frac{\eta_1 \delta}{\mu + l + d_H} \right) \left(1 + \frac{l}{\mu} + \frac{l \delta}{\mu (\mu + l + d_H)} \right) \right. \\ & \left. + \frac{2 \delta}{\mu + l + d_H} \right] + \left(\frac{2 \eta_2 l \delta}{\mu (\mu + l + d_H)} \right) \\ & \left(1 + \frac{\delta}{\mu + l + d_H} \right) + \frac{2l^2}{\mu^2} \left(1 + \frac{\delta}{\mu + l + d_H} \right)^2 \\ & + \left(\frac{l \delta}{\mu (\mu + l + d_H)} \right) \left(1 + \frac{\delta}{\mu + l + d_H} \right). \end{aligned}$$

If $v_1 > \bar{v}_1$, system (2) evinces a backward bifurcation (i.e., stable endemic equilibrium coexists with stable disease-free equilibrium) for $\mathcal{R}_E^H < 1$. On the other hand, if $v_1 < \bar{v}_1$ (i.e., $a < 0$) there exists a supercritical transcritical bifurcation at $\mathcal{R}_E^H = 1$. That is, the behavior of endemic equilibrium χ_H^* changes from unstable (for $\mathcal{R}_E^H < 1$) to stable (for $\mathcal{R}_E^H > 1$) while passing through $\mathcal{R}_E^H = 1$, associated with bifurcation parameter $\lambda_H^* = \lambda_H$. Thus, the proof is completed.

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