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Mathematical Modelling and Optimal Controls for Controlling Pneumonia-HIV Co-Infection

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Abstract:

Bacterial infections like Pneumonia have emerged as an important cause of morbidity and mortality in individuals infected with HIV giving rise to their co-infection. There are mathematical models that describe; the co-dynamics, treatment and protection mechanisms of Pneumonia and HIV infections. Therefore, this research is aimed at determining optimal control treatments through developing a deterministic mathematical model Pneumonia-HIV co-infection incorporating the use of anti-pneumonia and ART treatment interventions as controls. The uncontrolled pneumonia-HIV co-infection is presented with its analysis. Finally, the optimal control theory for Pneumonia-HIV co-infection model is derived analytically by applying the Pontryagin's Maximum Principle.

Keywords: Antiretroviral therapy, antiretroviral, pontryagin's maximum principle, basic reproduction number

1. Introduction

Human-Immunodeficiency Virus (HIV) is a virus that causes acquired-immunodeficiency syndrome (AIDS). It targets the immune system, specifically the CD4 cells or T cells of the infected individual weakening it and making an individual incapable of resisting attack from a wide range of infections (WHO, 2012). Currently, there is no cure or vaccination for HIV infected individuals but HIV Testing Services are being offered to people with HIV and AIDS to help in improving their quality of lives as well as controlling the spread. The general treatment of HIV positive people is with ARV and HAART drugs which can lower the viral load set point while prolonging the life of infected person (Ho, 1995). This treatment reduces the risk of HIV transmission. But before administering the ARV, an individual is first administered with prophylaxis for opportunistic, counseling and assessing the eligibility of ARV. Regardless of ART or HAART treatment, the infectiousness is not restored to susceptibility or reduced and hence an infected individual could still cause the infection. Research shows that there are millions approximately 34 million of individuals who have HIV and AIDS and died from the disease (UNAIDS, 2013; WHO, 2014). This is evident that further studies and researches on the disease are needed in order to reduce the number of infections.

On the other hand, Pneumonia is an infectious disease that is caused by microbes including bacteria, virus, and fungi among people with severely weakened immune systems or example those infected with HIV. It is classified as an airborne disease more generally a lung infection involving alveoli (air sacs). The common causes of pneumonia are Streptococcus Pneumoniae (bacterial), Pneumocystis Carinii Pneumonia (PCP) and histoplasma (fungal) and Haemophilus influenza type b and respiratory syncytial virus (RSV) (viral) (Gray and Zar, 2010; Greenwood, 2000). Currently, there is no specific effective prevention, treatment or vaccination of Pneumonia as its cause has several factors. Generally the disease can be vaccinated with pneumococcal conjugate vaccines or treated with antibiotics (cotrimoxazole prophylaxis, amoxillin, ampicillin and gentamycine dispersible tablets).

The co-infection due to Pneumonia and HIV results once the immune system has been impaired with HIV in which CD4 or T-cells are less than 200 cells per millimetre cubed giving a chance to Pneumonia infection. Research shows that the lungs are the principal target of HIV associated complications and persons with HIV infections are at an increased risk for a wide spectrum of opportunistic cases of pneumonia, neoplasms and pulmonary condition (WHO, 2012; Sogaard, *et al*, 2008). It has become a burden to the society disrupting the social-economical systems as efforts have been directed to curb these instances.

Several scholars have developed mathematical models which play a great role in proposing ways of controlling the transmission co-dynamics of HIV and pneumonia (Lutera *et al.*, 2018; J. Nthiiri *et al.*, 2015; K. Nthiiri *et al.*, 2016; Onyinge *et al.*, 2016).

These models did not incorporate the aspect of optimal controls. Therefore, this study aims at incorporating the aspect of optimal control theory (Pontryagin's Maximum Principle) through developing a mathematical model, analyzing and obtaining optimal treatment controls for co-infection of Pneumonia and HIV that that will be effective.

2. Model Formulation

We formulated a model for the co-infection of Pneumonia and HIV that subdivide the total population at time t , $N(t)$ into the following compartments:

$S(t)$: The number of individuals that are susceptible to both Pneumonia and HIV infections. This compartment is increased by recruits at the rate, Λ and it is increased from those individuals who have recovered from Pneumonia R_p after losing their temporary immunity at the rate δ . These individuals can get Pneumonia infection and move to the I_p at the rate λ_p and also can get HIV infection and move to the I_h class at the rate λ_h . λ_p and λ_h are given as:

$$\lambda_p = \frac{\phi_1(I_p(t) + \iota_1 I_{ph}(t) + \iota_2 A_{ph}(t))}{N(t)}, \lambda_h = \frac{\phi_2(I_h(t) + \beta_1 I_{ph}(t))}{N(t)} \quad (1)$$

where ϕ_1 is the effective transmission for Pneumonia and $\iota_2 > \iota_1$ are modification parameters which accounts for relative increased infectiousness with Pneumonia, ϕ_2 is the effective transmission rate for HIV and β_1 is a modification parameter for increased relative infectiousness of individuals with HIV.

$I_p(t)$: The number of individuals infected with Pneumonia. They are reduced by recoveries from Pneumonia at the rate ϵ_1 while others get infected with HIV and progress to I_{ph} class at the rate $\delta_1 \lambda_h$. Mortality induced by Pneumonia under I_p class occurs at the rate d_p . $R_p(t)$: The number of individuals who have recovered from Pneumonia infection. These individuals lose their temporary immunity and they get to susceptible class at the rate δ while others get infection from HIV at the rate λ_h and progress to I_h class.

$I_h(t)$: The number of individuals infected with HIV with no clinical symptoms of HIV and AIDS who progress to the class A_h by developing clinical symptoms of HIV and AIDS at the rate γ_1 while others get infected with Pneumonia and progress to I_{ph} class at the rate $\delta_2 \lambda_p$. $I_{ph}(t)$: The number of individuals co-infected with both Pneumonia and HIV with no clinical symptoms of HIV and AIDS. These individuals recover from Pneumonia at the rate ϵ_2 and move to I_h class, others develop HIV and AIDS symptoms and progress to A_{ph} class at the rate γ_2 while others die from Pneumonia induced mortality rate d_p .

$A_h(t)$: The number of individuals infected with HIV showing clinical symptoms of HIV and AIDS. These individuals seek medication from Pneumonia and they progress to A_{ph} class at a recovery rate ϵ_3 and are increased from A_{ph} class who get infection from Pneumonia at the rate $\delta_3 \lambda_p$ where $\delta_3 > \delta_2$. They incur induced death from both Pneumonia and AIDS at mortality rate d_{Ap} .

$A_{ph}(t)$: The number of individuals co-infected with both Pneumonia and HIV with clinical symptoms of HIV and AIDS. They incur AIDS induced death at the rate d_A . All classes are reduced by deaths from natural causes at the rate μ .

The resulting diagram from the definition of the model is given in Figure 1 and described mathematically by the system (2) as;

Figure 1: Flow Chart Diagram

$$\begin{aligned} \dot{S}(t) &= \Lambda + \delta R_p - (\lambda_p + \lambda_h + \mu)S \\ \dot{I}_p(t) &= \lambda_p S - (\epsilon_1 + \mu + d_p + \delta_1 \lambda_h)I_p \\ \dot{R}_p(t) &= \epsilon_1 I_p - (\mu + \delta + \lambda_h)R_p \\ \dot{I}_h(t) &= \lambda_h R_p + \epsilon_2 I_{ph} + \lambda_h S - (\mu + \gamma_1 + \delta_2 \lambda_p)I_h \\ \dot{I}_{ph}(t) &= \delta_1 \lambda_h I_p + \delta_2 \lambda_p I_h - (\gamma_2 + \epsilon_2 + \mu + d_p)I_{ph} \\ \dot{A}_{ph}(t) &= \gamma_2 I_{ph} + \delta_3 \lambda_p A_h - (\epsilon_3 + \mu + d_{Ap})A_{ph} \\ \dot{A}_h(t) &= \gamma_1 I_h + \epsilon_3 A_{ph} - (\delta_3 \lambda_p + \mu + d_A)A_h \end{aligned} \quad (2)$$

3. Model Analysis

Model (2) is analyzed in a feasible bounded region Π given as: $\Pi = \{(S(t), I_p(t), R_p(t),$

$I_h(t), I_{ph}(t), A_{ph}(t), A_h(t)) \in \mathbb{R}_+^7 : N(t) \leq \frac{\Lambda}{\mu}$. Since this model deals with the co-epidemic of the two diseases in a varying human population size we assumed all state variables and parameters to be non-negative for all time $t \geq 0$ which are bounded in the region Π . As $t \rightarrow \infty$, we have

$$0 \leq N(t) \leq \frac{\Lambda}{\mu} \quad (3)$$

Therefore the model is well posed and hence biologically meaningful.

4. Disease-Free Equilibrium Point (DFE)

DFE E^0 is a steady-state solution for which there is no disease or infection in the population (Olaniyi and Obabiyi, 2014). To obtain the disease-free equilibrium point we set each equation of the right handside of the model system (2) equal to zero and solve hand side the system. Since there are no infections in the human populations, we set $I_p(t) = R_p(t) = I_h(t) = I_{ph}(t) = A_{ph}(t) = A_h(t) = 0$ and obtain the disease-free equilibrium point E^0 of the model (2) given by: $E^0 = \{S(t), I_p(t), R_p(t), I_h(t), I_{ph}(t), A_{ph}(t), A_h(t)\} = (\frac{\Lambda}{\mu}, 0, 0, 0, 0, 0, 0)$

5. Endemic Equilibrium Point

Endemic equilibrium point E^* is a steady-state solution for which there exists a constant occurrence of diseases within the population (Olaniyi and Obabiyi, 2014). To obtain the endemic equilibrium point, we set each of the right handside of the model system (2) equal to zero and solve the resultant system using mathematics. The solutions are given as:

$$S^*(t) = \frac{-\Lambda - p}{\mu + \lambda_h + \lambda_p}, A_h^*(t) = \frac{\gamma_2 \epsilon_3}{Q_5} \left(\frac{Q_4}{Q_2} + \frac{3}{(\mu + d_p \epsilon_1 + \delta_1 \lambda_h) Q_2} \right) - \frac{\gamma_1 (-\mu - d_{Ap} - \epsilon_3)}{Q_5} \left(\frac{1(\Lambda - p)}{\delta_2 Q_1} \right), I_p^*(t) =$$

$$\frac{(\Lambda - \delta R_p) \lambda_p}{Q_1}, R_p^*(t) = \frac{-\Lambda - \delta R_p}{\mu + \lambda_h + \lambda_p} + \frac{\epsilon_1}{\lambda_h} \left(\frac{Q_4}{Q_2} + \frac{Q_3}{(\mu + d_p \epsilon_1 + \delta_1 \lambda_h) Q_2} \right) - \frac{(-\mu - \gamma_1 - \delta_2 \lambda_p)}{\lambda_p} \left(\frac{Q_4}{Q_2} + \frac{Q_3}{(\mu + d_p \epsilon_1 + \delta_1 \lambda_h) Q_2} \right),$$

$$I_h^*(t) = -\frac{\delta_1 (\Lambda - \delta R_p) \lambda_h}{\delta_2 Q_1} + \frac{(-\mu - d_p - \gamma_1 - \epsilon_2)}{\delta_2 \lambda_p} \left(\frac{Q_4}{Q_2} + \frac{Q_3}{(\mu + d_p \epsilon_1 + \delta_1 \lambda_h) Q_2} \right), I_{ph}^*(t) = \frac{-Q_4}{Q_2} - \frac{Q_3}{Q_2 (\mu + d_p \epsilon_1 + \delta_1 \lambda_h)},$$

$$A_{ph}^*(t) = \frac{\gamma_1}{\epsilon_3} \left(\frac{\delta_1 (\Lambda + \delta R_p) \lambda_h}{\delta_2 Q_1} - \frac{(-\mu - d_p - \gamma_1 - \epsilon_2)}{\delta_2 \lambda_p} \left(\frac{\gamma_2 \epsilon_3}{Q_5} \left(\frac{Q_4}{Q_2} + \frac{Q_3}{(\mu + d_p \epsilon_1 + \delta_1 \lambda_h) Q_2} \right) \right) \right) + \frac{\gamma_2 (-\mu - d_A - \delta_3 \lambda_p)}{Q_5} \left(\frac{\gamma_2 \epsilon_3}{Q_5} \right) -$$

$$\frac{\gamma_1 (-\mu - d_{Ap} - \epsilon_3)}{Q_5} \left(\frac{\delta_1 (\Lambda - \delta R_p) \lambda_h}{\delta_2 Q_1} \right)$$

where, $Q_1 = (\mu + d_p + \epsilon_1 + \delta_1 \lambda_h)(\mu + \lambda_h + \lambda_p)$, $Q_2 = (\mu + \lambda_h + \lambda_p)(-\delta_2 \epsilon_2(-\delta - \mu - \lambda_h) \lambda_p + (-\mu - d_p - \gamma_1 - \epsilon_2)(-\delta - \mu - \lambda_h)(-\gamma_1 - \mu - \delta_2 \lambda_p))$, $Q_3 = (\Lambda + \delta R_p) \lambda_p (\delta_2 \epsilon_1 \lambda_h \lambda_p + \delta_1 (-\delta - \mu - \lambda_h) \lambda_h (-\mu - \gamma_1 - \delta_2 \lambda_p))$, $Q_4 = \delta_2 (-\Lambda - \delta R_p)(\delta - \mu - \lambda_h) \lambda_h \lambda_p$, $Q_5 = \delta_3 \epsilon_3 \lambda_p - (-\mu - d_{Ap} - \epsilon_3)(-\mu - d_A - \delta_3 \lambda_p)$

6. The Basic Reproduction Number, R_0

We use next generation matrix approach to determine R_0 that measures the ability of a disease reproducing itself (Van den Driessche and Watmough, 2002). The two reproduction numbers at disease free equilibrium representing the two diseases are:

$$R_{0P} = \frac{\kappa_1}{(\epsilon_1 + \mu + d_p)}, R_{0H} = \frac{(\mu + d_A) \kappa_2 + \gamma_1 \kappa_2^2}{(\mu + \gamma_1)(\mu + d_A)} \quad (4)$$

$$R_0 = \max\{R_{0P}, R_{0H}\}$$

Such that;

7. Local Stability of the Disease Free Equilibrium Point

The Jacobian matrix of the model system (2) is used to evaluate the local stability of the system at E^0 . At DFE, the Jacobian matrix of equation (2) is

$$J_1 = \begin{pmatrix} -\mu & -\frac{\kappa_1 \Lambda}{\mu} & \delta & 0 & 0 & 0 & 0 \\ 0 & -(\epsilon_1 + \mu + d_p) & 0 & 0 & 0 & 0 & 0 \\ 0 & \epsilon_1 & -(\mu + \delta) & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & -(\mu + \gamma_1) & \epsilon_2 & 0 & 0 \\ 0 & 0 & 0 & 0 & B_1 & 0 & 0 \\ 0 & 0 & 0 & 0 & \gamma_2 & B_2 & 0 \\ 0 & 0 & 0 & \gamma_1 & 0 & \epsilon_3 & -(\mu + d_A) \end{pmatrix} \quad (5)$$

where, $B_1 = -(\gamma_2 + \epsilon_2 + \mu + d_p)$, $B_2 = -(\epsilon_3 + \mu + d_{Ap})$

From (5), the trace $\text{tra} J_1 = d_A - 4\mu - d_{Ap} - \gamma_1 - \gamma_2 + \delta_1 + 2\epsilon_1 - \epsilon_2 - \epsilon_3$ which is negative and determinant is given by $\det J_1(E^0) = -(-\mu - d_A) \mu (-\mu - \gamma_1) \epsilon_1 (-\mu - d_p - \delta_1 - \epsilon_2 - \gamma_2) (-\epsilon_3 - \mu - d_{Ap})$ which is positive if $(\mu - d_A) \mu (-\mu - \gamma_1) \epsilon_1 (-\mu - d_p - \delta_1 - \epsilon_2 - \gamma_2) (-\epsilon_3 - \mu - d_{Ap}) > 0$ (6)

Therefore, the disease free equilibrium point is asymptotically stable.

8. Local Stability of the Endemic Equilibrium Point

The Jacobian of equation (2) at endemic state E^* is

$$J_2 = \begin{pmatrix} b_1 & -\frac{\kappa_1 S^*}{N} & \delta & -\frac{\kappa_2 S^*}{N} & 0 & 0 & 0 \\ \frac{\kappa_1 I_p^*}{N} & b_2 & 0 & -\frac{\kappa_2 \delta_1 I_p^*}{N} & -\frac{\kappa_2 \delta_2 I_p^*}{N} & 0 & 0 \\ 0 & \epsilon_1 & b_3 & -\frac{\kappa_2 R_p^*}{N} & 0 & 0 & 0 \\ \frac{\kappa_2 I_h^*}{N} & -\frac{\delta_2 \kappa_1 I_h^*}{N} & \lambda_h^* & \kappa_1 \frac{R_p^* + S^*}{N} + b_4 & \epsilon_2 - \frac{\delta_2 \kappa_1 \iota_1 I_h^*}{N} & 0 & 0 \\ 0 & \delta_1 \lambda_h^* + \frac{\delta_2 \kappa_1 I_h^*}{N} & 0 & \delta_2 \lambda_p^* + \frac{\delta_1 \kappa_2 I_p^*}{N} & b_5 + b_6 & 0 & 0 \\ 0 & \frac{\delta_3 \kappa_1 A_h^*}{N} & 0 & 0 & \gamma_2 & b_7 + \frac{\delta_3 \iota_2 \kappa_1 A_h^*}{N} & \frac{\delta_3 \iota_2 \kappa_1 A_{ph}^*}{N} \\ 0 & \frac{\delta_3 \kappa_1 A_h^*}{N} & 0 & \gamma_1 & 0 & \epsilon_3 - \frac{\iota_2 \kappa_1 \delta_3 A_h^*}{N} & b_8 \end{pmatrix} \quad (7)$$

where $b_1 = -(\frac{\kappa_1 I_p^*}{N} + \frac{\kappa_1 I_h^*}{N} + \mu)$, $b_2 = -(\epsilon_1 + \mu + d_p + \delta_1 \lambda_h^*)$, $b_3 = -(\mu + \delta + \lambda_h^*)$, $b_4 = -(\mu + \gamma_1 + \delta_2 \lambda_p^*)$, $b_5 = -(\gamma_2 + \epsilon_2 + \mu + d_p)$, $b_6 = \frac{\delta_2 \kappa_1 \iota_1 I_h^* + \delta_2 \kappa_2 I_p^*}{N}$, $b_7 = -(\epsilon_3 + \mu + d_{Ap})$, $b_8 = -(\delta_3 \lambda_p^* + \mu + d_A)$, $\lambda_p^* = \frac{\kappa_1 (I_p^* + \iota_1 I_{ph}^*)}{N}$, $\lambda_p^{**} = \frac{\kappa_1 (I_p^* + \iota_2 A_{ph}^*)}{N}$, $\lambda_h^* = \frac{\kappa_2 (I_h^* + \iota_1 I_{ph}^*)}{N}$

From (7), the trace of $J_2 = -d_A - \delta - 7\mu - d_{Ap} - 2d_p - \gamma_1 - \gamma_2 - \epsilon_1 - \epsilon_2 - \epsilon_3 - \frac{\kappa_1 I_h^*}{N} - \frac{\kappa_1 I_p^*}{N} + \frac{\kappa_1 (S^* + R_p^*)}{N} + \frac{\delta_3 \iota_2 \kappa_1 A_h^*}{N} + \frac{\delta_2 \kappa_1 \iota_1 I_h^* + \delta_2 \kappa_2 I_p^*}{N} - \lambda_h - \delta_1 \lambda_h - \delta_2 \lambda_p - \delta_3 \lambda_p$ Which is negative if

$$\frac{\kappa_1 (S^* + R_p^*)}{N} + \frac{\delta_3 \iota_2 \kappa_1 A_h^*}{N} + \frac{\delta_2 \kappa_1 \iota_1 I_h^* + \delta_2 \kappa_2 I_p^*}{N} - \lambda_h - \delta_1 \lambda_h - \delta_2 \lambda_p - \delta_3 \lambda_p < 0 \quad (8)$$

And determinant
 $\det J_2(E^*)$

$$\begin{aligned} &= -\frac{1}{N} A_{ph} \delta_3 \iota_2 \kappa_1 \left(\epsilon_3 - \frac{A_h \delta_3 \iota_2 \kappa_1}{N} \right) - \frac{1}{N} \left(\epsilon_2 - \frac{i_h \delta_1 \iota_1 \kappa_1}{N} \right) \left(\frac{\delta I_p \epsilon_1 \kappa_1}{N} + (-\delta - \mu - \lambda_h) \left(\frac{S I_p \kappa_1^2}{N^2} \right) \right) \\ &+ \left(\epsilon_2 - \frac{I_h \delta_1 \epsilon_1 \iota_1 \kappa_1}{N} \right) + \left(\frac{\delta I_p \epsilon_1 \kappa_1}{N} + (-\mu - \delta - \lambda_h) \right) + (-\mu - \epsilon_1 - \gamma_2 - d_p) + \frac{S I_p \kappa_2 \kappa_1}{N^2} \\ &+ \left(\frac{S I_p \kappa_1^2}{N} + (-\mu - \frac{I_h \kappa_1}{N} - \frac{I_p \kappa_1}{N}) ((-\mu - \epsilon_1 - \delta_1 \lambda_h - d_p)) \right) \\ &+ \left(\frac{I_h \delta_2 \iota_1 \kappa_1 + I_p \delta_2 \iota_2 \kappa_1}{N} \right) \left(\frac{R_p^2 \kappa_2^2 \delta}{N} \left(\frac{I_h I_p \delta_2 \kappa_2^2}{N^2} + \frac{\mu I_h \kappa_2^2}{N} + \frac{I_h d_p \kappa_2^2}{N} + \frac{I_h \epsilon_1 \kappa_2^2}{N} + \frac{I_h \delta_1 \kappa_2^2 \lambda_h}{N} \right) \right) \\ &+ \left(\frac{(\mu - \delta - \lambda_h)(\delta_2 \kappa_1)}{N} \right) \left(\frac{\mu I_p \delta_1 \kappa_2}{N^2} + \frac{S I_p \kappa_2 \kappa_1}{N^2} + \frac{I_h I_p \delta_1 \kappa_1 \kappa_2^2}{N^2} + \frac{I_p^2 \delta_1 \kappa_1 \kappa_2^2}{N^2} \right) + \left(\frac{(\mu - \delta - \lambda_h)(\delta_2 \kappa_1)}{N} \right) \left(\frac{\mu I_p \delta_1 \kappa_2}{N^2} \right) \\ &+ \frac{I_h I_p \delta_1 \kappa_1 \kappa_2^2}{N^2} + \frac{I_p^2 \delta_1 \kappa_1 \kappa_2^2}{N^2} + \left(\frac{(\mu - d_p - \epsilon_1 - \delta_1 \lambda_h)(I_p \kappa_1)}{N} \right) \left(\frac{S \kappa_1}{N} + \frac{\delta R_p \kappa_1}{N} + S \frac{I_h \kappa_2^2}{N^2} + \frac{S I_p \kappa_2 \kappa_1}{N^2} - \delta \delta_2 \lambda_p \right) - \\ &(\epsilon_2 - \frac{\delta_1 I_h \iota_1 \kappa_1}{N}) \left(\frac{\delta I_p R_p \kappa_1 \kappa_2}{N^2} - \frac{\mu \delta_1 I_p \kappa_1}{N} - S \frac{I_p \kappa_2 \kappa_1}{N^2} - \frac{S I_h I_p \delta_1 \kappa_2 \kappa_1}{N^2} - \frac{S I_h^2 \delta_1 \kappa_2 \kappa_1}{N^2} \right) (-\delta - \mu - \lambda_h) - \\ &\left(\frac{-d_A - \mu - \delta_1 \lambda_p}{N} \right) (\delta_2 I_p \kappa_1) + (\epsilon_2 - \frac{\delta_1 I_h \iota_1 \kappa_1}{N}) \left(\frac{\delta I_p \epsilon_1 \kappa_1}{N} \right) + \delta \frac{I_p R_p \kappa_2 \kappa_1}{N^2} - \frac{S I_h I_p \delta_1 \kappa_2 \kappa_1}{N^2} - \frac{I_p I_h \delta_1 \kappa_2 \kappa_1}{N^2} \\ &+ \frac{I_p^2 \delta_1 \kappa_2 \kappa_1}{N^2} + \frac{I_p \delta_1 \mu \kappa_2 \kappa_1}{N^2} + \frac{\delta I_p R_p \kappa_1 \kappa_2}{N^2} - \left(\frac{\mu \delta_1 I_p \kappa_1}{N} - \frac{S I_p \kappa_2 \kappa_1}{N^2} - \frac{S I_h I_p \delta_1 \kappa_2 \kappa_1}{N^2} - \frac{S I_h^2 \delta_1 \kappa_2 \kappa_1}{N^2} \right) (-\delta - \mu - \lambda_h) \end{aligned}$$

$\det J_2 > 0$ if

$$(-\delta - \mu - \lambda_h) > 0 \quad (9)$$

Therefore, the endemic equilibrium point is asymptotically stable provided the inequalities (8) and (9) hold otherwise it is unstable.

9. Extension of the Model into Optimal Control

We incorporate the two controls interventions, u_1 : Pneumonia treatment and u_2 : HIV treatment using the concepts of optimal control theory (Pontryagin *et al.*, 1962) to get extended model as:

$$\begin{aligned} \dot{S}(t) &= \Lambda + \delta R_p - (\lambda_p + \lambda_h + \mu)S \\ \dot{I}_p(t) &= \lambda_p S - (\epsilon_1 + u_1 + \mu + d_p + \delta_1 \lambda_h) I_p \\ \dot{R}_p(t) &= (\epsilon_1 + u_1) I_p - (\mu + \delta + \lambda_h) R_p \\ \dot{I}_h(t) &= \lambda_h R_p + (\epsilon_2 + u_1) I_{ph} + \lambda S - (\mu + \gamma_1(1 - u_2) + \delta_2 \lambda_p) I_h \\ \dot{I}_{ph}(t) &= \delta_1 \lambda_h I_p + \delta_2 \lambda_p I_h - (\gamma_2(1 - u_2) + (\epsilon_2 + u_1) + \mu + d_p) I_{ph} \\ \dot{A}_{ph}(t) &= \gamma_2(1 - u_2) I_{ph} + \delta_3 \lambda_p A_h - ((\epsilon_3 + u_1) + \mu + d_{Ap}) A_{ph} \\ \dot{A}_h(t) &= \gamma_1(1 - u_2) I_h + (\epsilon_3 + u_1) A_{ph} - (\delta_3 \lambda_p + \mu + d_A) A_h \end{aligned} \quad (10)$$

To study the optimal control levels, the control set U is considered to be Lebesgue measurable and it is defined as: $U = \{(u_1(t), u_2(t)) : 0 \leq u_1 < 1, 0 \leq u_2 < 1, 0 \leq t \leq T\}$.

The proposed objective function J given by:

$$J = \int_{t_0}^T \left(k_1 I_p + k_2 I_h + k_3 I_{ph} + k_4 A_{ph} + k_5 A_h + \frac{1}{2} w_1 u_1^2 + \frac{1}{2} w_2 u_2^2 \right) dt \quad (11)$$

where $k_1, k_2, k_3, k_4, k_5, w_1$ and w_2 are positive balancing coefficients (weights) which regularize the optimal control.

We need to obtain optimal controls (u_1^*, u_2^*) such that:

$$J(u_1^*, u_2^*) = \min \{ J(u_1(t), u_2(t)) : u_1, u_2 \in U \} \quad (12)$$

subject to the dynamical system above (10) and the control set U .

10. Existence of the Optimal Control Problem

The basic framework of an optimal control is to prove the existence of the optimal control to characterize it for optimality. The conditions that are to be met include;

- The integrand of the objective functional is convex on U .
- The set of controls and corresponding state variables (admissible pairs) A is not empty. (iii) The control set U is compact.
- There exists non-negative constants τ_1 and τ_2 and $\phi > 1$ such that the integrand of the objective function is bounded by $\tau_1 + \tau_2 (|u_1|^2 + |u_2|^2)^{\phi/2}$
- Each right-hand side of the state system is continuous and bounded above by a linear function in the state and control variables.
- We note that all the above conditions are met, there exists unique optimal controls which are unique as a result of the Lipschitz property of the state system with respect to the state variables and a priori boundedness of the state solutions (Fleming and Rishel, 2012).

11. Characterization of Optimal Controls

We consider the necessary conditions to determine the optimal controls given in (12) with constraint model (10) could be obtained using the Pontryagin Maximum Principle.

$$H(S, I_p, R_p, I_h, I_{ph}, A_{ph}, A_h, u_1, u_2, \zeta_1, \zeta_2, \zeta_3, \zeta_4, \zeta_5, \zeta_6, \zeta_7) = k_1 I_p + k_2 I_h + k_3 I_{ph} + k_4 A_{ph} + k_5 A_h + \frac{1}{2} w_1 u_1^2 + \frac{1}{2} w_2 u_2^2 + \sum_{i=1}^7 \zeta_i l_i \quad (13)$$

where l_i are the i th state variable equations on the right-hand side of equation (10) and $\zeta_1, \dots, \zeta_7$ are the adjoint satisfying the following theorem.

Theorem: For optimal controls (u_1^, u_2^*) and solution set $\{(S(t), I_p(t), R_p(t), I_h(t), I_{ph}(t), A_{ph}(t), A_h(t))\}$ of the corresponding state system that minimizes the objective function J over U , there exist adjoint variables $\zeta_1, \dots, \zeta_7$ such that:*

$$\begin{aligned} \dot{\zeta}_1(t) &= \zeta_1(\lambda_p + \lambda_h + \mu) - \zeta_2 \lambda_p - \zeta_4 \lambda_p \\ \dot{\zeta}_2(t) &= -k_1 + \zeta_2(\epsilon_1 + u_1 + \mu + d_p + \delta_1 \lambda_h) - \zeta_3(\epsilon_1 + u_1) - \zeta_5 \delta_1 \lambda_h \\ \dot{\zeta}_3(t) &= -\zeta_1 \delta + \zeta_3(\mu + \delta + \lambda_h) - \zeta_4 \lambda_h \\ \dot{\zeta}_4(t) &= -k_2 + \zeta_4(\mu + \gamma_1(1 - u_2) + \delta_2 \lambda_p) - \zeta_5 \delta_2 \lambda_p - \zeta_7 \gamma_1(1 - u_2) \\ \dot{\zeta}_5(t) &= -k_3 - \zeta_4(\epsilon_2 + u_1) + \zeta_5(\gamma_2(1 - u_2) + \epsilon_2 + u_1 + \mu + d_p) - \zeta_6 \gamma_2(1 - u_2) \\ \dot{\zeta}_6(t) &= -k_4 + \zeta_6(\epsilon_3 + u_1 + \mu + d_{Ap}) - \zeta_7(\epsilon_3 + u_1) \\ \dot{\zeta}_7(t) &= -k_5 + \zeta_7(\delta_3 \lambda_p + \mu + d_A) - \zeta_6 \delta_3 \lambda_p \end{aligned} \quad (14)$$

with transversality conditions; $\zeta_1(T) = \zeta_2(T) = \zeta_3(T) = \zeta_4(T) = \zeta_5(T) = \zeta_6(T) = \zeta_7(T) = 0$.

Furthermore, the optimal controls are represented by;

$$\begin{aligned} u_1^* &= \max\{0, \min(1, \frac{(\zeta_2 - \zeta_3)I_p + (\zeta_4 - \zeta_5)I_{ph} + (\zeta_6 - \zeta_7)A_{ph}}{w_1})\} \\ u_2^* &= \max\{0, \min(1, \frac{(\zeta_4 - \zeta_7)I_h + (\zeta_5 - \zeta_6)I_{ph}}{w_2})\} \end{aligned} \quad (15)$$

Proof: From the relation in

$$\dot{\zeta}_i(t) = -\frac{\partial H}{\partial x^i} \quad (16)$$

where $i = 1, \dots, 7$, we get:

$$\begin{aligned}
\dot{\zeta}_1(t) &= \zeta_1(\lambda_p + \lambda_h + \mu) - \zeta_2\lambda_p - \zeta_4\lambda_h \\
\dot{\zeta}_2(t) &= -k_1 + \zeta_2(\epsilon_1 + u_1 + \mu + d_p + \delta_1\lambda_h) - \zeta_3(\epsilon_1 + u_1) - \zeta_5\delta_1\lambda_h \\
\dot{\zeta}_3(t) &= -\zeta_1\delta + \zeta_3(\mu + \delta + \lambda_h) - \zeta_4\lambda_h \\
\dot{\zeta}_4(t) &= -k_2 + \zeta_4(\mu + \gamma_1(1 - u_2) + \delta_2\lambda) - \zeta_5\delta_2\lambda_p - \zeta_7\gamma_1(1 - u_2) \\
\dot{\zeta}_5(t) &= -k_3 - \zeta_4(\epsilon_2 + u_1) + \zeta_5(\gamma_2(1 - u_2) + \epsilon_2 + u_1 + \mu + d_p) - \zeta_6\gamma_2(1 - u_2) \\
\dot{\zeta}_6(t) &= -k_4 + \zeta_6(\epsilon_3 + u_1 + \mu + d_{Ap}) - \zeta_7(\epsilon_3 + u_1) \\
\dot{\zeta}_7(t) &= -k_5 + \zeta_7(\delta_3\lambda_p + \mu + d_A) - \zeta_6\delta_3\lambda_p
\end{aligned} \tag{17}$$

With transversality conditions; $\zeta_1(T) = \zeta_2(T) = \zeta_3(T) = \zeta_4(T) = \zeta_5(T) = \zeta_6(T) = \zeta_7(T) = 0$ $\square$

We use optimality condition $\frac{\partial H}{\partial u_i} = 0$ to obtain the time varying optimal controls as below:

$$\frac{\partial H}{\partial u_1} = u_1 w_1 + (\zeta_3 - \zeta_2)I_p + (\zeta_5 - \zeta_4)I_{ph} + (\zeta_7 - \zeta_6)A_{ph} = 0$$

$$\frac{\partial H}{\partial u_2} = u_2 w_2 + (\zeta_4 - \zeta_7)I_h + (\zeta_5 - \zeta_6)I_{ph} = 0 \tag{18}$$

we obtain the following after re-arranging and simplifying;

$$u_1^* = \frac{(\zeta_2 - \zeta_3)I_p + (\zeta_4 - \zeta_5)I_{ph} + (\zeta_6 - \zeta_7)A_{ph}}{w_1}$$

$$u_2^* = \frac{(\zeta_4 - \zeta_7)I_h + (\zeta_5 - \zeta_6)I_{ph}}{w_2} \tag{19}$$

Applying boundary conditions of each control we get;

$$u_1^* = \max\{0, \min(1, \frac{(\zeta_2 - \zeta_3)I_p + (\zeta_4 - \zeta_5)I_{ph} + (\zeta_6 - \zeta_7)A_{ph}}{w_1})\}$$

$$u_2^* = \max\{0, \min(1, \frac{(\zeta_4 - \zeta_7)I_h + (\zeta_5 - \zeta_6)I_{ph}}{w_2})\} \tag{20}$$

12. Conclusion

The main purpose of this research work was to determine optimal controls for controlling the spread of Pneumonia and HIV co-infection. This study therefore developed a deterministic model incorporating treatment controls for the co-infection which was shown to be positively invariant as well as bounded. The Pontryagin's maximum principle was used to derive and analyze the necessary conditions for the optimal controls (use of anti-Pneumonia and use of ARVs).

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