Effect of therapy on the dynamics of HIV model with adaptive immune response and two saturated rates

Karam Allali (with S. Harroudi and Y. Tabit)

Laboratory of Mathematics and Applications, University of Casablanca, Morocco

17th International Symposium on Mathematical and Computational Biology

Institute of Numerical Mathematics, Russian Academy of Sciences, Moscow, Russia, - 30th October - 03rd November 2017



2 The model

Ositivity and Boundedness

Analyse of the model

- Stability of the disease-free equilibrium
- Stabiliy of the infection steady states

5 Conclusion

6 References

Introduction

- Human Immunodeficiency virus (HIV) is a pathogen which causes the well known Acquired Immunodeficiency Syndrom (AIDS).
- The HIV dynamics involving the density uninfected cells, the density of the infected cells, the density of HIV virus and the amount of CTL cells have been widely studied starting from the work by Nowak and Bangham (1996)¹ and two years later by De Boer et al. (1998)².
- The dynamic of HIV including CTL cells and two saturated rates is studied by Tabit et al. (2014)³

¹Nowak M.A.; Bangham C.R.M. Population dynamics of immune responses to persistent viruses. Science 173 1996, 272, 7479.

²De Boer, Rob J., Perelson. Alan S. Target cell limited and immune control models of HIV infection: a 200 comparison, Journal of theoretical Biology 190 (1998) 201214.

The model

The dynamics of HIV infection with CTL, antibody responses and therapy that we consider is given by the following nonlinear system of differential equations

$$\begin{cases} \frac{dT}{dt} = s - dT - \frac{(1 - \eta)\beta VT}{1 + aV} + \rho I, \\ \frac{dI}{dt} = \frac{(1 - \eta)\beta VT}{1 + aV} - (\delta + \rho)I - \rho IZ, \\ \frac{dV}{dt} = (1 - \epsilon)N\delta I - \mu V - qVW, \end{cases}$$
(1)
$$\frac{dW}{dt} = gVW - hW, \\ \frac{dZ}{dt} = \frac{cIZ}{1 + \alpha I} - bZ. \end{cases}$$

With the initial conditions $T(0) = T_0$, $I(0) = I_0$, $V(0) = V_0$, $Z(0) = Z_0$ and $W(0) = W_0$.

The solutions of the problem (1) exist. Moreover, they are bounded, nonnegative and verify:

i)
$$T_1(t) \leq T_1(0) + \frac{s}{\delta_1}$$
,
ii) $V(t) \leq V(0) + \frac{(1-\epsilon)N\delta}{\mu} ||I||_{\infty}$,
iii) $W(t) \leq W(0) + \frac{g}{q} [\max(1; 2-\frac{\mu}{h})V(0) + (\frac{(1-\epsilon)N\delta}{\mu} + \frac{(1-\epsilon)N\delta}{h}) ||I||_{\infty}]$,
iv) $Z(t) \leq Z(0) + \frac{c}{p} [\max(1; 2-\frac{d}{b})T(0) + I(0) + \max(\frac{s}{b}; \frac{s}{d}) + \max(0; 1-\frac{\delta}{b}) ||I||_{\infty}]$,
where $T_1(t) = T(t) + I(t)$ and $\delta_1 = \min(d; \delta)$.

э

・ロン ・四 と ・ ヨ と ・ ヨ と

Elements of the Proof

Proof (Elements)

• We have
$$\dot{T}_1 = s - dT - \delta I - p I Z$$
, thus

$$T_1(t) \leq T_1(0)e^{-\delta_1 t} + rac{s}{\delta_1}(1-e^{-\delta_1 t})$$

• From $\dot{V} = (1 - \epsilon)N\delta I - \mu V - qVW$, we have

$$V(t) \leq V(0)e^{-\mu t} + (1-\epsilon)N\delta\int_0^t I(\xi)e^{(\xi-t)\mu}d\xi$$

See

$$\dot{W} + hW = gVW = \frac{g}{q} \left((1 - \epsilon)N\delta I - (\dot{V} + \mu V) \right)$$

• From
$$\dot{Z} = \frac{cIZ}{1+\alpha I} - bZ$$
 we have
 $\dot{Z} + bZ \le cIZ = \frac{c}{p}[s - (\dot{T} + dT) - (\dot{I} + \delta I)]$

6/29

The basic reproduction number of the system is given by

$$R_0 = \frac{(1-\theta)N\delta s}{d\mu(\delta+\rho)}.$$

$$(1-\theta) = (1-\eta) \times (1-\epsilon)$$

There is an infection-free equilibrium

$$E_f = \left(\frac{s}{d}, 0, 0, 0, 0\right)$$

corresponding to the maximal level of healthy CD4+ T-cells.

- The disease-free equilibrium, E_f, is locally asymptotically stable for R₀ < 1.</p>
- 2 The disease-free equilibrium, E_f , is unstable for $R_0 > 1$.

Elements of the Proof

Proof (Elements) At the disease-free equilibrium, E_f , the Jacobian matrix is given as follows:

$$J_{E_{f}} = \begin{pmatrix} -d & \rho & -\frac{(1-\eta)\beta s}{d} & 0 & 0\\ 0 & -(\delta+\rho) & \frac{(1-\eta)\beta s}{d} & 0 & 0\\ 0 & (1-\epsilon)N\delta & -\mu & 0 & 0\\ 0 & 0 & 0 & -h & 0\\ 0 & 0 & 0 & 0 & -b \end{pmatrix}$$
(3)

The characteristic polynomial of J_{E_f} is

$$P_{E_f}(\xi) = (\xi + d)(\xi + b)(\xi + h)[\xi^2 + (\delta + \rho + \mu)\xi + (\delta + \rho)\mu(1 - R_0)],$$

One of the eigenvalues is

$$\xi = \frac{-(\delta + \rho + \mu) + \sqrt{(\delta + \rho + \mu)^2 - 4(\delta + \rho)\mu(1 - R_0)}}{2},$$

Stability of the disease-free equilibrium



Figure 1: Behavior of the infection during the time which correspond to the stability of the free-equilibrium E_{f} . s = 5, $\beta = 0.000024$, d = 0.02, $\delta = 0.5$, p = 0.001, N = 500, $\mu = 3$, $\rho = 0.01$, a = 0.001, $\alpha = 0.001$, c = 0.03, b = 0.2, q = 0.5, $g = 10^{-11}$, h = 0.1, $\eta = 0.4$ and $\epsilon = 0.55$.

(a)

11/29

The infection steady states: $E_1 = (T_1, I_1, V_1, 0, 0)$, where

$$T_{1} = \frac{s}{d} \left[\frac{a(1-\epsilon)Ns + \mu}{a(1-\epsilon)Ns + \mu R_{0}} \right],$$

$$I_{1} = \frac{s}{\delta} \left[\frac{\mu(R_{0}-1)}{a(1-\epsilon)Ns + \mu R_{0}} \right],$$

$$V_{1} = \frac{(1-\epsilon)Ns(R_{0}-1)}{a(1-\epsilon)Ns + \mu R_{0}},$$

 $E_2 = (T_2, I_2, V_2, W_2, 0)$, where

$$T_{2} = \frac{(\rho + \delta)(g + ah)s}{d(\rho + \delta)(g + ah) + (1 - \eta)\beta\delta h},$$

$$I_{2} = \frac{(1 - \eta)\beta hs}{d(\rho + \delta)(g + ah) + (1 - \eta)\beta\delta h},$$

$$V_{2} = \frac{h}{g},$$

$$W_{2} = \frac{\mu}{q} [\frac{(1 - \theta)N\delta\beta gs}{\mu[d(\rho + \delta)(g + ah) + (1 - \eta)\beta\delta h]} - 1],$$

 $E_3 = (T_3, I_3, V_3, 0, Z_3)$, where

$$\begin{split} I_3 &= \frac{b}{c - \alpha b}, \\ T_3 &= \frac{(a(1 - \epsilon)N\delta\rho)I_3^2 + (a(1 - \epsilon)Ns\delta + \mu\rho)I_3 + \mu s)}{(1 - \epsilon)N\delta(ad + (1 - \eta)\beta)I_3 + \mu d)}, \\ V_3 &= \frac{(1 - \epsilon)N\delta I_3}{\mu}, \\ Z_3 &= \frac{-(1 - \epsilon)N\delta[ad\rho + \delta(ad + (1 - \eta)\beta)]I_3}{p((1 - \epsilon)N\delta(ad + (1 - \eta)\beta)I_3 + \mu d)} \\ &+ \frac{((1 - \theta)\beta Ns\delta - d\mu(\rho + \delta))}{p((1 - \epsilon)N\delta(ad + (1 - \eta)\beta)I_3 + \mu d)}, \end{split}$$

and $E_4 = (T_4, I_4, V_4, W_4, Z_4)$, where

$$I_{4} = \frac{b}{c - \alpha b},$$

$$V_{4} = \frac{h}{g},$$

$$T_{4} = \frac{(s + \rho I_{4})(1 + aV_{4})}{d(1 + aV_{4}) + (1 - \eta)\beta V_{4}},$$

$$W_{4} = \frac{1}{q}(\frac{(1 - \epsilon)N\delta I_{4}}{V_{4}} - 1),$$

$$Z_{4} = \frac{1}{p}(\frac{s}{I_{4}} - \frac{dT_{4}}{I_{4}} - \delta).$$

< □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □

In order to study the local stability of the points E_1 , E_2 , E_3 and E_4 , we first define the following numbers:

$$D_0^W = \frac{(1-\epsilon)gNs}{h\mu}, \ \widetilde{D_0^W} = D_0^W \frac{\mu R_0}{(a(1-\epsilon)Ns + \mu R_0)}, \ H_0^W = \frac{1}{\frac{1}{R_0} + \frac{1}{\widetilde{D_0^W}}},$$

$$D_0^Z = \frac{cs}{b\delta}, \ \widetilde{D_0^Z} = D_0^Z \frac{\mu \delta R_0}{(a(1-\epsilon)Ns + \mu R_0) + \alpha \mu s(R_0 - 1))}, \ H_0^Z = \frac{1}{\frac{1}{R_0} + \frac{1}{\widetilde{D_0^Z}}},$$

and

$$H_0^{W,Z} = \frac{D_0^Z R_0}{D_0^W (1 + \frac{ah}{g}) + R_0 (1 + \frac{\alpha s^2}{\delta})}.$$

・ロ ・ ・ 一 ・ ・ 注 ト ・ 注 ・ う へ で
16 / 29

- If $R_0 < 1$, then the point E_1 does not exist.
- **2** If $R_0 = 1$, then $E_1 = E_f$.
- If $R_0 > 1$, then E_1 is locally asymptotically stable for $H_0^W < 1$, and $H_0^Z < 1$; however it is unstable for $H_0^W > 1$ or $H_0^Z > 1$.

Proof (Elements)

It easy to see that if $R_0 < 1$, then the point E_1 does not exist and if $R_0 = 1$ the two points E_1 and E_f coincide. If $R_0 > 1$, the Jacobian matrix at E_1 is given by



then, its characteristic equation is

$$(\xi + h - gV_1)(\xi + b - \frac{cI_1}{1 + \alpha I_1})(\xi^3 + a_1\xi^2 + a_2\xi + a_3) = 0,$$

where

$$\begin{aligned} a_1 &= d + \delta + \mu + \rho + \frac{(1 - \eta)\beta V_1}{1 + aV_1}, \\ a_2 &= (\delta + \mu + \rho)d + (\mu + \delta)\frac{(1 - \eta)\beta V_1}{1 + aV_1} + \mu(\delta + \rho) - \frac{(1 - \theta)N\delta\beta T_1}{(1 + aV_1)^2}, \\ a_3 &= \mu d(\delta + \rho) + \frac{\mu\delta(1 - \eta)\beta V_1}{1 + aV_1} - \frac{(1 - \theta)N\delta\beta T_1 d}{(1 + aV_1)^2}, \end{aligned}$$

We have $gV_1 - h = \frac{h\widetilde{D_0^W}(H_0^W - 1)}{H_0^W}$ and $\frac{cl_1}{1 + \alpha l_1} - b = \frac{b\widetilde{D_0^Z}(H_0^Z - 1)}{H_0^Z}.$ Checking the negativity by Routh-Hurwitz Theorem.



Figure 2: Behavior of the infection during the time which correspond to the stability of the endemic-equilibrium point E_1 . s = 5, $\beta = 0.00024$, d = 0.02, $\delta = 0.5$, p = 0.001, N = 1200, $\mu = 3$, $\rho = 0.01$, a = 0.001, $\alpha = 0.001$, c = 0.03, b = 0.2, q = 0.5, $g = 10^{-11}$, h = 0.1, $\eta = 0.1$ and $\epsilon = 0.2$.

- If $H_0^W < 1$, then the point E_2 does not exist.
- **2** If $H_0^W = 1$ then $E_2 = E_1$.
- If $H_0^W > 1$ then E_2 is locally asymptotically stable for $H_0^{W,Z} < 1$ and unstable for $H_0^{W,Z} > 1$.



Figure 3: Behavior of the infection during the time which correspond to the stability of the endemic-equilibrium E_2 , s = 10, $\beta = 0.000024$, d = 0.02, $\delta = 0.5$, p = 0.001, N = 1200, $\mu = 3$, $\rho = 0.01$, a = 0.001, $\alpha = 0.001$, c = 0.03, b = 0.2, q = 0.001, $g = 10^{-4}$, h = 0.01, $\eta = 0.55$ and $\epsilon = 0.45$.

<□▶ < □▶ < □▶ < 三▶ < 三▶ < 三 りへぐ 22/29

• If
$$\alpha > \frac{c}{b}$$
 or $H_0^Z < 1$, then the point E_3 does not exist and $E_3 = E_2$ when $H_0^Z = 1$.

2 If
$$\alpha < \frac{c}{b}$$
, $H_0^Z > 1$ and $D_0^W < (1 - \frac{\alpha b}{c})D_0^Z$, then E_3 is locally asymptotically stable.

3 If
$$\alpha < \frac{c}{b}$$
, $H_0^Z > 1$ and $D_0^W > (1 - \frac{\alpha b}{c})D_0^Z$, then E_3 is unstable.



Figure 4: Behavior of the infection during the time which correspond to the stability of the endemic-equilibrium E_3 , s = 15, $\beta = 0.000024$, d = 0.02, $\delta = 0.5$, p = 0.001, N = 1200, $\mu = 3$, $\rho = 0.01$, a = 0.001, $\alpha = 0.001$, c = 0.03, b = 0.2, q = 0.5, $g = 10^{-11}$, h = 0.1, $\eta = 0.02$ and $\epsilon = 0.07$.



Figure 5: Behavior of the infection during the time which correspond to the stability of the endemic-equilibrium E_4 , s = 10, $\beta = 0.000024$, d = 0.02, $\delta = 0.5$, p = 0.001, N = 1200, $\mu = 3$, $\rho = 0.01$, a = 0.001, $\alpha = 0.001$, c = 0.03, b = 0.2, q = 0.5, $g = 10^{-4}$, h = 0.1, $\eta = 0.05$ and $\epsilon = 0.2$.

- The local stability of the disease-free equilibrium depends on the basic reproduction number R_0 .
- The local stability of the infection steady states depends on the basic reproduction number R_0 , the CTL immune response reproduction number D_0^Z and the antibody immune response reproduction number D_0^W .
- In the presence of therapy, an increases of the uninfected cells is observed.
- The results of this work confirm that the therapy may control the viral replication and reduce the infection.

References I

- Nowak M.A.; Bangham C.R.M. Population dynamics of immune responses to persistent viruses. Science 173 1996, 272, 7479.
- De Boer, Rob J., Perelson. Alan S. Target cell limited and immune control models of HIV infection: a 200 comparison, Journal of theoretical Biology 190 (1998) 201214.
- Y. Tabit, A. Meskaf, K. Allali. Mathematical analysis of HIV model with two saturated rates, CTL and antibody responses. *World Journal of Modelling and Simulation*, 2016 12(2): 137–146.
- Y. Tabit, K. Hattaf, N. Yousfi. Dynamics of an HIV pathogenesis model with CTL immune response and two saturated rates. *World Journal of Modelling and Simulation*, 2014 **10**(3): 215–223.
- K. Allali, Y. Tabit, S. Harroudi. On HIV model with adaptive immune response, two saturated rates and therapy .Math. Model. Nat. Phenom. Vol. 12, No. 5, 2017, pp. 114.

Thank you for your attention

イロト イポト イヨト イヨト 一日

29/29