

# GLOBAL MATHEMATICAL ANALYSIS OF AN HIV INFECTION MODEL WITH FULL LOGISTIC GROWTH AND SATURATION INCIDENCE

Liancheng Wang\* and Xiaoqin P. Wu<sup>†‡</sup>

**Abstract.** In this research, we study an HIV infection mathematical model with a full logistic growth and a saturation incidence. Both local and global mathematical analysis are carried out. By identifying a basic reproduction number  $R_0$ , the existence and stability of the uninfected steady state  $P_0$  and the unique infected steady state  $P^*$  are established in terms of  $R_0$ . We show that if  $R_0 \leq 1$ ;  $P_0$  is the only equilibrium in the feasible region, it is globally asymptotically stable. Therefore, no HIV infection persists and infected T cells and HIV virus are cleared over time. However, If  $R_0 > 1$ , then the unique infected steady state  $P^*$  emerges.  $P_0$  becomes unstable. We show that the system is uniformly persistent and HIV infection persists. The unique infected steady state  $P^*$  is globally asymptotically stable under some conditions.

**Keywords.** HIV infection model, global analysis, steady states, uniform persistence, stability.

## 1 Introduction

The most notorious disease of AIDS/HIV has been studied intensively for the last several decades. Researchers are working diligently to find a cure for the disease from many different aspects including mathematical viewpoint. Different mathematical models that describe the interaction of HIV virus and T cells have been developed and studied, see for instance [1-11] and references therein. Typically, to describe the HIV infection, we partition the total T cell population into healthy (uninfected) T cells and infected T cells and study the dynamics between healthy T cells, infected T cells, and HIV free virus with their concentrations represented by  $T(t)$ ,  $T^*(t)$  and  $V(t)$  at time  $t$ , respectively. Then a basic mathematical model describing the HIV infection can be written in the follow-

ing form,

$$\begin{aligned}\dot{T} &= s - \alpha T - kVT, \\ \dot{T}^* &= kVT - \beta T^*, \\ \dot{V} &= N\beta T^* - \epsilon V,\end{aligned}\tag{1}$$

where all parameters  $s, \alpha, \beta, \epsilon, k, N$  are nonnegative. The human body produces T cells and it is assumed that T cells are produced at a constant rate  $s$  and that newly produced T cells are healthy and susceptible to HIV virus. Parameters  $\alpha, \beta$ , and  $\epsilon$  are the per-capita death rates of the healthy T cells, infected T cells, and the virus particles, respectively. It is natural to assume that  $\alpha \leq \beta$ , i.e. the infected T cells have a greater death rate than healthy cells because of the infectivity of the T cells. The infection is through a direct contact between virus and healthy T cells. The incidence is described by a simple mass-action term  $kVT$ , where  $k > 0$  is the contact rate between virus and healthy T cells. Infected T cells produce virus and each infected T cell produces  $N$  virus during its lifetime.

It is well known now that T cells proliferate once stimulated by antigen or mitogen. Even though this proliferation process is still not well understood, a logistic proliferation form has been widely used in literature. Perelson and Nelson [3] and Leenheer and Smith [8] studied model (1) with a simplified logistic form,  $rT(1 - T/T_{max})$ , where  $T_{max}$  is the capacity T cell population. With this simplified logistic form, the system is competitive and a complete mathematical analysis is carried out by Leenheer and Smith [8] using the theory for three dimensional competitive systems. Recently, Wang and Li [9] proposed and studied model (1) with a full logistic proliferation term,  $rT(1 - (T + T^*)/T_{max})$ , and Wang and Ellermeier [10] proposed and studied a similar model that utilized the full logistic form in both the first and second equations assuming that both the healthy and infected T cells proliferate. With the full logistic proliferation form, the system is no longer competitive and the mathematical analysis is carried out using an approach developed by Li and Muldowney [12]. More recently, Wang [11] studied system (1) with a Holling type-II incidence form. Local and global stability analysis are carried out.

In this paper, we will focus on system (1) with a full-

\*Liancheng Wang is with Department of Mathematics and Statistics, Kennesaw State University, USA. E-mails: l-wang5@kennesaw.edu

<sup>†</sup>Xiaoqin P. Wu is with Department of Mathematics, Computer and Information Sciences, Mississippi Valley State University, USA. E-mail: xpaul.wu@yahoo.com

<sup>‡</sup>Manuscript received January 1, 2011; revised April 1, 2011.