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Effects of co

Infection of human immunodeficient healthy CD4⁺ T-cells in a well-mixed model is considered to be a transcription plus protease inhibition model. This model is used to explain the steady states in HIV growth. A numerical method (VIM), is used to solve the model. The results are rapidly convergent successive approximations are obtained without any initial guess. The physical behavior of the problem is illustrated. The effects of proposed

Keywords: HIV infection; CD4⁺ T-cells; cacy.

Mathematics Subject Classification

1. Introduction

Human immunodeficiency virus (HIV) is a retrovirus. Its hallmark is the reverse transcription of its RNA into DNA. The gp120 protein of HIV on the surface is a 55-kDa protein found on the surface of T-helper cells. It has a helper function in the immune system. It is found in T-helper cells/macrophages and dendritic cells. It undergoes a conformational change