

**Stability Analysis of an SVIR Epidemic Model with Nonlinear Incidence Rate**

**การวิเคราะห์เสถียรภาพของแบบจำลองการระบาด SVIR ที่มีอัตราอุบัติการณ์ที่ไม่เป็นเชิงเส้น**

Aphirak Boonpikum (อภิรักษ์ บุญพิคำ)\* Dr.Wirawan Chinvirasit (ดร.วิราวรรณ ชินวิริยสิทธิ์)\*\*

**ABSTRACT**

This paper, an SVIR epidemic model with nonlinear incidence rate is investigated. The stability of the model depends on the basic reproduction number. The global asymptotically stable of the disease-free is proved by constructing a Lyapunov function and LaSalle’s invariant set theorem. An unique endemic equilibrium exists and is locally asymptotically stable whenever  $R_0 > 1$ . Numerical simulation is illustrated to support the analytical results and to study the effect of nonlinear incidence rate on the SVIR model.

**บทคัดย่อ**

งานวิจัยนี้ได้สำรวจแบบจำลองการระบาด SVIR ที่มีอัตราอุบัติการณ์ที่ไม่เป็นเชิงเส้น ซึ่งเสถียรภาพของแบบจำลองนี้ขึ้นอยู่กับค่าระดับการติดเชื้อของโรค การสร้างฟังก์ชันไลออปโนฟและทฤษฎีเซตไม่แปรผันของลาซาล ได้นำไปพิสูจน์เสถียรภาพเชิงเส้นกำกับวงกว้างของจุดสมดุลภายใต้สภาวะไร้โรค และจุดสมดุลภายใต้สภาวะการแพร่ระบาดมีเสถียรภาพเฉพาะที่เชิงเส้นกำกับเมื่อ  $R_0 > 1$  นอกจากนี้งานวิจัยได้แสดงผลลัพธ์เชิงตัวเลขเพื่อใช้สนับสนุนผลลัพธ์เชิงทฤษฎีและศึกษาผลกระทบของอัตราอุบัติการณ์ไม่เชิงเส้นในแบบจำลอง SVIR

**Key Words:** SVIR , Vaccination, Lyapunov function

**คำสำคัญ:** แบบจำลองการระบาด SVIR การให้วัคซีน ฟังก์ชันไลออปโนฟ

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\*Student, Master of Science, Program in Applied Mathematics, Department of Mathematics, Faculty of Science, King Mongkut’s University of Technology Thonburi

\*\* Associate Professor, Department of Mathematics, Faculty of Science, King Mongkut’s University of Technology Thonburi

**1. Introduction**

Mathematical models in epidemiology or epidemic model are used to understand epidemiological phenomena, to predict the infectious disease progress and to prevent public health for controlling the diseases. Most epidemic models descend from the classical *SIR* (Susceptible-infectious-recovered) model of Kermack and Mckendric established in 1927 (W.O. Kermarck and A.G. Mckendrick, 1927). In fact, vaccines are extremely important and have been proved to be the most effective and cost-efficient method of preventing infectious diseases such as measles, polio, diphtheria, tetanus, pertussis, and tuberculosis. In recent years, many researchers have discussed the *SIR* model allowing vaccination, that is, the *SVIR* model.

Kribs-Zaleta C.M. and Velasco-Hernández J.X. (2000) presented a *SVIS* epidemic model and found that this model exhibited the phenomenon of backward bifurcation. Furthermore, the model is applied for studying the vaccination of disease pertussis and tuberculosis. Liu X., Takeuchi Y. and Iwami S. (2008) established two *SVIR* model to describe continuous vaccination strategy and pulse vaccination strategy, respectively. However, in modeling disease epidemics take place in ecological system, the incidence rate plays a key factor in the transmission of infectious diseases. It has been suggested by several authors that the disease transmission process may have a nonlinear incidence rate. This allows one to include behavioral changes and prevent unbounded contact rates. A particular example of such an incidence rate is given by  $\beta I(1 + \nu I^{k-1})S$ , where  $\beta > 0$ ,  $\nu > 0$  and  $k > 0$ . This incidence with  $\nu$  near one, represents saturation or multiple exposures before infection. van der Driessche P. and Watmough J. (2000, 2003) introduced this incident into *SIS* epidemic model where  $\beta > 0$ ,  $\nu > 0$  and  $k > 0$ .

They found that *SIS* epidemic model with this incidence rate exhibited backward bifurcation. Alexander M.E. and Moghadas S.M. (2004) analyzed an *SIV* model with a generalized nonlinear incidence rate  $\beta I(1 + \nu I^q)S$ , where  $\beta > 0$ ,  $\nu \geq 0$  and  $0 < q \leq 1$ . The results showed that bistability and various Hopfbifurcation occurred. Jin Y., Wang W. and Xiao S. (2007) studied the backward bifurcation, the Hopf bifurcation and Bogdanov-Takens bifurcation by analyzing a *SIRS* model with nonlinear incident rate in the form  $\beta I(1 + \nu I^{k-1})S$ , where  $\beta > 0$ ,  $\nu > 0$  and  $k = 2$ . Zhou X. and Cui J. (2011) investigated the *SEIV* model with nonlinear incidence rate of the form  $\beta I(1 + \nu I^{q-1})S$  with  $\beta > 0$ ,  $\nu > 0$  and  $q = 2$ .

The aim of this paper is to investigate the effect of nonlinear incidence rate  $\beta SI(1 + \nu I)$ ,  $\beta > 0$  and  $0 < \nu \leq 1$  on the dynamic of a *SVIR* epidemic model proposed by Liu X., Takeuchi Y. and Iwami S. (2008). To this end, we consider the following differential equations

$$\begin{aligned}
 \frac{dS}{dt} &= \mu - \mu S - \beta SI(1 + \nu I) - \alpha S, \\
 \frac{dV}{dt} &= \alpha S - \beta_1 VI - \gamma_1 V - \mu V, \\
 \frac{dI}{dt} &= \beta SI(1 + \nu I) + \beta_1 VI - \gamma I - \mu I, \\
 \frac{dR}{dt} &= \gamma_1 I + \gamma I - \mu R,
 \end{aligned}
 \tag{1}$$

where the state variables  $S, V, I$  and  $R$  denote the densities (or fractions) of susceptible, vaccinated, infected and recovered individuals, respectively. The parameters  $\mu$  is the natural birth rate which is assumed to be equal natural death rate;  $\beta$  is the transmission rate;  $\alpha$  is the rate at which susceptible individuals who are moved into the vaccination process;  $\beta_1$  is the possibility of infection with a disease transmission rate while contracting with

infected individuals;  $\gamma$  is the recovery rate of infected individuals and  $\gamma_1$  is the average rate for individuals to obtain immunity and move into recovered population, respectively. It is assumed that  $\beta_1$  is less than  $\beta$  because the vaccinated individuals may have some partial immunity during the process or they may recognize the transmission characters of the disease and hence decrease the effective contacts with infected individuals. Adding all equations in (1) gives

$$\frac{dN}{dt} = \frac{dS}{dt} + \frac{dV}{dt} + \frac{dI}{dt} + \frac{dR}{dt} = \mu - \mu N \quad (2)$$

which has the following implication: the three-dimension simplex

$$\Gamma = \left\{ (S, V, I, R) \in \mathbb{R}_+^4 : S + V + I + R = 1 \right\}$$

is positively invariant. On the simplex  $\Gamma$ , we have  $R(t) = 1 - S(t) - V(t) - I(t)$ , thus, the dynamics of system (1) is studied by analyzing the following three-dimensional system:

$$\begin{aligned} \frac{dS}{dt} &= \mu - \mu S - \beta SI(1 + \nu I) - \alpha S, \\ \frac{dV}{dt} &= \alpha S - \beta_1 VI - \gamma_1 V - \mu V, \\ \frac{dI}{dt} &= \beta SI(1 + \nu I) + \beta_1 VI - \gamma I - \mu I, \end{aligned} \quad (3)$$

The dynamical behavior of (1) on  $\Gamma$  is equivalent to that of (3). Therefore, the system (3) is studied in the feasible region

$$\Gamma = \left\{ (S, V, I) \in \square_+^3 : 0 \leq S, V, I \leq 1, 0 \leq S + V + I \leq 1 \right\} \quad (4)$$

where  $\square_+^3$  denotes the non-negative cone of  $\square^3$  including its lower-dimensional faces.

The paper is organized as follows: In section 2, the model is analyzed the existence of equilibriums in the model. The SVIR model is conserved to analyze the stability analysis which consists of the disease-free equilibrium and endemic equilibria. The

Lyapunovfunction is analyzed for proving the global stability under the condition and the locally asymptotically stable is analyzed both equilibriums in Section 3. The last section, Numerical simulations are performed in the last section.

## 2. Existence of Equilibriums

In the absence of infection, by setting the right-hand side of system (3) to zero and solving the obtained results, the model has a disease-free equilibrium  $E_0$ ,

$$E_0 = (S_0, V_0, I_0) = \left( \frac{\mu}{\mu + \alpha}, \frac{\alpha\mu}{(\mu + \alpha)(\mu + \gamma_1)}, 0 \right) \quad (5)$$

and an endemic equilibrium  $E^*$ , with

$$\begin{aligned} S^* &= \frac{\mu}{\mu + \alpha + \beta I^* (1 + \nu I^*)}, \\ V^* &= \frac{\alpha S^*}{\mu + \gamma_1 + \beta_1 I^*}, \end{aligned} \quad (6)$$

where  $I^*$  is positive root of cubic equation

$$h(I^*) = wI^{*3} + xI^{*2} + yI^* + z = 0, \quad (7)$$

with

$$\begin{aligned} w &= \nu\beta\beta_1(\mu + \gamma) > 0, \\ x &= \nu\beta(\mu + \gamma)(\mu + \gamma_1) + \gamma\beta\beta_1 \\ &\quad + (1 - \nu)\mu\beta\beta_1 > 0, \\ y &= (\mu + \gamma)(\beta(\mu + \gamma_1) + \beta_1(\mu + \alpha)) \\ &\quad - \nu\beta\mu(\mu + \gamma_1) - \mu\beta\beta_1, \\ z &= (\mu + \gamma_1)(\mu + \alpha)(\mu + \gamma)(1 - R_0), \\ R_0 &= \frac{\beta\mu}{(\mu + \alpha)(\mu + \gamma)} + \frac{\beta_1\alpha\mu}{(\mu + \alpha)(\mu + \gamma)(\mu + \gamma_1)}. \end{aligned} \quad (8)$$

Here,  $R_0$  is called the basic reproduction number of system (3). Thus, the following Theorem is established.

**Theorem 1.** The system (3) has a unique endemic equilibrium  $E^*$  for  $R_0 > 1$ , and has no endemic equilibrium for  $R_0 \leq 1$ .

**Proof.** Since  $0 < v \leq 1$ . It is clear that, from (8)  $w, x > 0$  and Then, the roots of (7) depend on the sign of  $y$  and  $z$ . Now, two cases of  $R_0$  are considered as follows:

Case 1:  $R_0 \leq 1$ , then

$$\begin{aligned} y &= (\mu + \gamma)(\beta(\mu + \gamma_1) + \beta_1(\mu + \alpha)) - v\beta\mu(\mu + \gamma_1) \\ &\quad - v\beta\mu(\mu + \gamma_1) - \mu\beta\beta_1 \\ &> \beta\mu(\mu + \gamma_1) + \beta_1(\mu + \gamma)(\mu + \alpha) - \beta\mu(\mu + \gamma_1) \\ &\quad - \beta\mu(\mu + \gamma_1) - \mu\beta\beta_1 \\ &> \beta_1(\mu + \gamma)(\mu + \alpha) \left( 1 - \frac{\mu\beta}{(\mu + \gamma)(\mu + \alpha)} \right. \\ &\quad \left. - \frac{\alpha\mu\beta_1}{(\mu + \gamma_1)(\mu + \alpha)(\mu + \gamma)} \right) \\ &= \beta_1(\mu + \gamma)(\mu + \alpha)(1 - R_0) > 0, \\ z &= (\mu + \gamma_1)(\mu + \alpha)(\mu + \gamma)(1 - R_0) > 0. \end{aligned}$$

Thus, if  $R_0 \leq 1$ , we have  $w > 0$ ,  $x > 0$ ,  $y > 0$  and  $z > 0$ . By the Descartes's rules of signs, there is no endemic equilibrium.

Case 2:  $R_0 > 1$ , then

$$\begin{aligned} y &= \beta(\mu + \gamma)(\mu + \gamma_1) + \beta_1(\mu + \gamma)(\mu + \alpha) \\ &\quad - v\beta\mu(\mu + \gamma_1) - \mu\beta\beta_1 \\ &= \beta(\mu + \gamma)(\mu + \gamma_1) \left( 1 - \frac{v\mu}{(\mu + \gamma)} \right) \\ &\quad + \beta_1(\mu + \gamma)(\mu + \alpha) \left( 1 - \frac{\mu\beta}{(\mu + \gamma)(\mu + \alpha)} \right) \\ &> \beta\mu(\mu + \gamma_1) \left( 1 - \frac{v\mu}{(\mu + \gamma)} \right) \\ &\quad + \beta_1\mu\alpha \left( 1 - \frac{\mu\beta}{(\mu + \gamma)(\mu + \alpha)} \right) \end{aligned}$$

$$\begin{aligned} y &> (\mu + \gamma)(\mu + \alpha)(\mu + \gamma_1) \left( \frac{\beta\mu}{(\mu + \gamma)(\mu + \alpha)} \right. \\ &\quad \times \left( 1 - \frac{v\mu}{(\mu + \gamma)} \right) + \frac{\beta_1\mu\alpha}{(\mu + \gamma)(\mu + \alpha)(\mu + \gamma_1)} \\ &\quad \left. \times \left( 1 - \frac{\mu\beta}{(\mu + \gamma)(\mu + \alpha)} \right) \right) \end{aligned}$$

Since

$$R_0 = \frac{\mu\beta}{(\mu + \gamma)(\mu + \alpha)} + \frac{\beta_1\mu\alpha}{(\mu + \gamma)(\mu + \alpha)(\mu + \gamma_1)} > 1,$$

we get

$$\frac{\beta_1\mu\alpha}{(\mu + \gamma)(\mu + \alpha)(\mu + \gamma_1)} > 1 - \frac{\mu\beta}{(\mu + \gamma)(\mu + \alpha)}.$$

It follows that

$$\begin{aligned} y &> (\mu + \gamma)(\mu + \alpha)(\mu + \gamma_1) \left( \frac{\beta\mu}{(\mu + \gamma)(\mu + \alpha)} \left( 1 - \frac{v\mu}{(\mu + \gamma)} \right) \right. \\ &\quad \left. + \left( 1 - \frac{\mu\beta}{(\mu + \gamma)(\mu + \alpha)} \right) \left( 1 - \frac{\mu\beta}{(\mu + \gamma)(\mu + \alpha)} \right) \right) \\ &= (\mu + \gamma)(\mu + \alpha)(\mu + \gamma_1) \left( \frac{\beta\mu}{(\mu + \gamma)(\mu + \alpha)} \left( 1 - \frac{v\mu}{(\mu + \gamma)} \right) \right. \\ &\quad \left. + \left( 1 - \frac{\mu\beta}{(\mu + \gamma)(\mu + \alpha)} \right)^2 \right) \end{aligned}$$

$$\text{and } z = (\mu + \gamma_1)(\mu + \alpha)(\mu + \gamma)(1 - R_0) < 0.$$

Thus,  $w > 0$ ,  $x > 0$ ,  $y > 0$  and  $z < 0$  if  $R_0 > 1$ . By the Descartes's rules of signs, there is an unique positive endemic equilibrium. This completes the proof.

### 3. Stability of equilibrium

**Theorem 2.** The disease-free equilibrium  $E_0$  is locally asymptotically stable if  $R_0 < 1$  and unstable if  $R_0 > 1$ .

**Proof.** The Jacobian of system (3) evaluated at  $E_0$  is

$$J(E_0) = \begin{bmatrix} -(\mu + \alpha) & 0 & \frac{-\beta\mu}{\mu + \alpha} \\ \alpha & -(\mu + \gamma_1) & \frac{-\beta_1\mu\alpha}{(\mu + \alpha)(\mu + \gamma_1)} \\ 0 & 0 & (\mu + \gamma)(R_0 - 1) \end{bmatrix} \quad (9)$$

with eigenvalues

$$\lambda_1 = -(\mu + \alpha) < 0, \quad \lambda_2 = -(\mu + \gamma_1) < 0 \quad (10)$$

$$\text{and } \lambda_3 = (\mu + \gamma)(R_0 - 1).$$

It follows that all eigenvalues (10) are negative real part. Therefore, the disease-free equilibrium  $E_0$  is locally asymptotically stable.

**Theorem 3.** If  $R_{01} \leq 1$ , then the disease-free equilibrium  $E_0$  of system (3) is globally asymptotically stable, where

$$R_{01} = \frac{(\beta\mu(\mu + \gamma_1) + \beta_1\alpha\mu)(1 + \nu)}{(\mu + \alpha)(\mu + \gamma)(\mu + \gamma_1)}.$$

**Proof.** Consider a Lyapunov function

$$L(S, V, I) = S - S_0 - S_0 \ln\left(\frac{S}{S_0}\right) + V - V_0 - V_0 \ln\left(\frac{V}{V_0}\right) + I,$$

its derivative along the solution of the system (3) is

$$\begin{aligned} \frac{dL}{dt} &= \frac{dS}{dt} + \frac{dV}{dt} + \frac{dI}{dt} - \frac{S_0}{S} \cdot \frac{dS}{dt} - \frac{V_0}{V} \cdot \frac{dV}{dt} \\ &= \mu - \mu S - \mu V - \mu I - \gamma_1 V - \gamma I - \frac{\mu S_0}{S} \\ &\quad + \mu S_0 + \beta S_0 I (1 + \nu I) + \alpha S_0 - \frac{\alpha S V_0}{V} \\ &\quad + \beta_1 V_0 I + \gamma_1 V_0 + \mu V_0 \\ &= -\mu S - \frac{\mu \alpha S_0}{V} + 2\mu S_0 + 3\alpha S_0 - \frac{S_0}{S} (\mu S_0 + \alpha S_0) \\ &\quad - \alpha \frac{V_0 S}{V} - (\mu + \gamma - \beta S_0 (1 + \nu I) - \beta_1 V_0) I \\ &\leq -\mu S_0 \left( \frac{S}{S_0} + \frac{S_0}{S} - 2 \right) - \alpha S_0 \left( \frac{V}{V_0} + \frac{S V_0}{S_0 V} + \frac{S_0}{S} - 3 \right) \\ &\quad - (\mu + \gamma - \beta S_0 (1 + \nu I) - \beta_1 V_0 (1 + \nu I)) I \\ &= -\mu S_0 \left( \frac{S}{S_0} + \frac{S_0}{S} - 2 \right) - \alpha S_0 \left( \frac{V}{V_0} + \frac{S V_0}{S_0 V} + \frac{S_0}{S} - 3 \right) \\ &\quad - (\mu + \gamma - (\beta S_0 + \beta_1 V_0) (1 + \nu I)) I \\ &= -\mu S_0 \left( \frac{S}{S_0} + \frac{S_0}{S} - 2 \right) - \alpha S_0 \left( \frac{V}{V_0} + \frac{S V_0}{S_0 V} + \frac{S_0}{S} - 3 \right) \\ &\quad - (\mu + \gamma) (1 - R_{01}) I \end{aligned}$$

Since all the model parameters are non-negative, it follows that  $L' \leq 0$  for all  $S, V, I > 0$  if  $R_{01} \leq 1$  and  $L' = 0$  if and only if  $S = S_0, V = V_0$  and  $I = 0$ . Now using LaSalle's invariance principle (J.P. LaSalle, 1967) of Lyapunov method, the limit set of each solution is contained in the largest compact invariant set  $\{(S, V, I) \in \Gamma : L'(S, V, I) = 0\}$ , which is the singleton  $\{E_0\}$ . Therefore, every solution that starts in  $\Gamma$  approaches  $E_0$  as  $t \rightarrow \infty$ . This completes the proof.

**Theorem 4.** If  $R_0 > 1$ , then the endemic equilibrium  $E^*$  of system (3) is locally asymptotically stable in  $\Gamma$ , provided  $a_1 a_2 - a_3 > 0$ , where  $a_1, a_2$  and  $a_3$  are presented in the proof.

**Proof.** The local stability for  $E^*$  is governed by the matrix is

$$J(E^*) = \begin{bmatrix} -\frac{\mu}{S^*} & 0 & -\beta S^* (1 + 2\nu I^*) \\ \alpha & -\frac{\alpha S^*}{V^*} & -\beta_1 V^* \\ \beta I^* (1 + \nu I^*) & \beta_1 I^* & \nu \beta S^* I^* \end{bmatrix}$$

The characteristic equation of  $J(E^*)$  is

$$\lambda^3 + a_1 \lambda^2 + a_2 \lambda + a_3 = 0,$$

where

$$a_1 = \frac{\mu}{S^*} + \frac{\alpha S^*}{V^*} - \nu \beta S^* I^*,$$

$$\begin{aligned} a_2 &= \frac{\alpha \mu}{V^*} + \beta_1^2 V^* I^* + \beta^2 S^* I^* (1 + 2\nu I^*) \\ &\quad + \nu \beta^2 S^* I^* (1 + 2\nu I^*) - \nu \beta S^* I^* \left( \frac{\mu}{S^*} + \frac{\alpha S^*}{V^*} \right), \end{aligned}$$

$$\begin{aligned} a_3 &= \alpha \beta \beta_1 S^* I^* (1 + 2\nu I^*) + \beta^2 \alpha \frac{S^{*2} I^*}{V^*} (1 + \nu I^*) (1 + 2\nu I^*) \\ &\quad + \beta_1^2 \mu \frac{V^* I^*}{S^*} > 0 \end{aligned}$$

It is seen that

$$\begin{aligned} a_1 &= \frac{\mu}{S^*} + \frac{\alpha S^*}{V^*} - v\beta S^* I^* \\ &= \beta I^* (1 + vI^*) + \beta_1 I^* + (\mu + \gamma_1) + (\mu + \alpha) - v\beta S^* I^* \\ &\geq \beta I^* + v\beta I^{*2} + \beta_1 I^* + (\mu + \gamma_1) + (\mu + \alpha) - \beta I^* \\ &= v\beta I^{*2} + \beta_1 I^* + (\mu + \gamma_1) + (\mu + \alpha) \\ &\geq v\beta I^{*2} + \beta_1 I^* + (\mu + \gamma_1) + (\mu + \alpha) \\ &= \beta_1 I^* (1 + vI^*) + (\mu + \gamma_1) + (\mu + \alpha) > 0. \end{aligned}$$

We also get

$$\begin{aligned} a_1 a_2 - a_3 &= \left( \frac{\mu}{S^*} + \frac{\alpha S^*}{V^*} - v\beta S^* I^* \right) \left( \frac{\alpha \mu}{V^*} + \beta_1^2 V^* I^* \right. \\ &\quad \left. + \beta^2 S^* I^* (1 + v)(1 + 2vI^*) - v\beta S^* I^* \left( \frac{\mu}{S^*} + \frac{\alpha S^*}{V^*} \right) \right) \\ &\quad - \left( \beta \alpha (\mu + \gamma) \frac{S^* I^* (1 + 2vI^*)}{V^*} + \beta_1^2 \mu \frac{V^* I^*}{S^*} \right) \\ &= \left( \frac{\mu}{S^*} + \frac{\alpha S^*}{V^*} - v\beta S^* I^* \right) \left( \frac{\alpha \mu}{V^*} + \beta_1^2 V^* I^* \right. \\ &\quad \left. + \beta^2 S^* I^* (1 + v)(1 + 2vI^*) - v\beta S^* I^* \left( \frac{\mu}{S^*} + \frac{\alpha S^*}{V^*} \right) \right) \\ &\quad \times \left( \frac{\mu}{S^*} + \frac{\alpha S^*}{V^*} - v\beta S^* I^* \right) - \beta \alpha (\mu + \gamma) \frac{S^* I^* (1 + 2vI^*)}{V^*} \\ &\quad - \beta_1^2 \mu \frac{V^* I^*}{S^*}. \end{aligned}$$

According to the Routh-Hurwitz criterion, the endemic equilibrium  $E^*$  is locally asymptotically stable in  $\Gamma$  if  $a_1 a_2 - a_3 > 0$ . This completes the proof.

#### 4. Numerical simulations

To investigate the dynamical behavior of the model, the system (3) is integrated numerically with parameters values:  $\mu = 0.01$ ,  $\gamma = 1/5$ ,  $\gamma_1 = 1/7$ ,  $\alpha = 1$  and  $v = 0.5$  with various  $\beta$  and  $\beta_1$ .

With above parameter values  $\beta = 4$  and  $\beta_1 = 1.25$ , the value of the basic reproduction number is  $R_0 = 0.51297$ , which yields  $R_{01} = 0.76946 < 1$ . Therefore, the system (3) has a disease-free equilibrium  $E_0(0.00990, 0.06520, 0)$  and the solutions of the

model (3) in the Figure 1 converge monotonically to  $E_0$ , as guaranteed by Theorem 1 and 2.

With above parameter values  $\beta = 8$  and  $\beta_1 = 6.5$ , the value of the basic reproduction number is  $R_0 = 2.38205 > 1$  and the system has a endemic equilibrium  $E^*(0.00793, 0.02240, 0.03094)$ . Figure 2 shows all solutions of the model (3) converge to  $E^*$  of the model (3) is locally asymptotically stable that is the disease persists.

For parameter values used:  $\mu = 0.8$ ,  $\beta = 3.96$ ,  $\beta_1 = 2.0$ ,  $\gamma = 0.5$ ,  $\gamma_1 = 0.4$ ,  $\alpha = 1.0$  for each value  $v$ ;  $v = 0, 0.5, 1.0$  and given by  $R_0 = 1.9236467 > 1$ . Thus, the system (3) has a unique equilibrium

$$E^*(0.2580750, 0.1390115, 0.3282510),$$

$$E^*(0.2272379, 0.1174843, 0.367099),$$

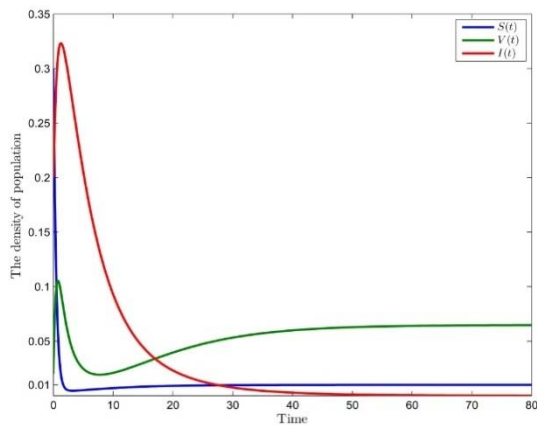
$$\text{and } E^*(0.1984950, 0.0990708, 0.4017839),$$

where  $v = 0, 0.5$  and  $v = 1.0$ , respectively. Figure 3 shows that the solution are locally asymptotically stable as guaranteed by Theorem 1 and 4. Moreover, it is found, see Figure 3, that even  $R_0$  does not depend on  $v$  but the density of infected population increases as  $v$  increases. This verifies that non-linear incidence rate has effected to the density of infectious populations.

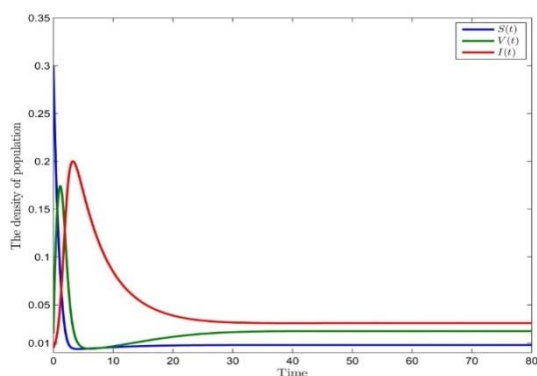
#### 5. Conclusion

In this paper we study the global stability of the *SVIR* model with nonlinear incidence rate  $\beta SI(1 + vI)$ , where  $\beta > 0$  and  $0 < v \leq 1$ . Our result show that the model has the reproduction number is similar to the model, which is proposed by Lie et al. (2008). (i.e., the reproduction number of the study system (3) does not depend on the parameter  $v$  in the nonlinear incidence function). The following results were obtained:

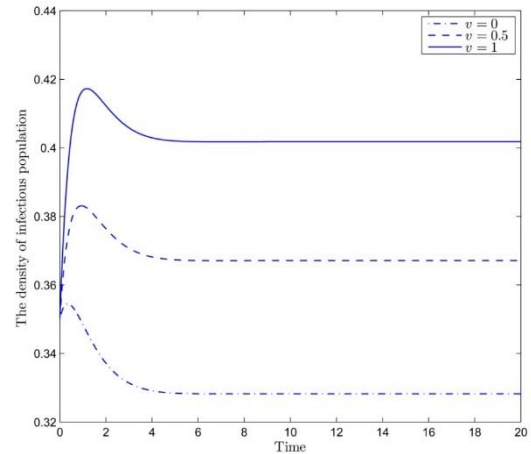
- (i) The model has a locally and globally asymptotically stable whenever  $R_{01} \leq 1$ .
- (ii) A unique endemic of the model has a locally asymptotically stable whenever the basic reproduction number is above the unity.
- (iii) In the numerical simulation of this model, the parameters  $\nu$  and  $\beta_1$  are effect to the infectious population and an endemic equilibrium that those parameters measure the effects of sociological, psychological or other mechanisms disease.



**Figure 1** Time series plots for the system (3) with parameter values used:  
 $\mu = 0.01, \beta = 4, \beta_1 = 1.25,$   
 $\gamma = 1/5, \gamma_1 = 1/7$  and  $\nu = 0.5$ .



**Figure 2** Time series plots for the system (3) with parameter values used:  
 $\mu = 0.01, \beta = 8, \beta_1 = 6.5,$   
 $\gamma = 1/5, \gamma_1 = 1/7$  and  $\nu = 0.5$ .



**Figure 3** Timeseries plots for the density of infectious individuals  $I(t)$  of the system (3) with parameter values used  $\mu = 0.8, \beta = 3.96, \beta_1 = 2.0, \gamma = 0.5, \gamma_1 = 0.4, \alpha = 1.0$  to compare the different parameters  $\nu$ ;  $\nu = 0, 0.5$  and  $\nu = 1.0$ , respectively.

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