

Bifurcation Analysis in Simple SIS Epidemic Model Involving Immigrations with Treatment

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Abstract: In this paper, local and Hopf bifurcation for an SIS epidemic model with treatment is investigated. Through theoretical analysis, we show the disease free equilibrium point has a transcritical bifurcation and the endemic equilibrium point has a saddle-node bifurcation. Also this model has a Hopf bifurcation near all equilibrium points. Applying the normal form theory and the center manifold argument, we derive the explicit formulas determining the properties of the bifurcation periodic solutions. In addition, we also study numerical simulations are also included.

Keywords: SIS epidemic model, Immigrations, Treatment, local and hopf bifurcation

1 Introduction

In recent decades, periodic oscillations (Bifurcation) received great attention due to they have been observed in the incidence of many infectious diseases, including measles, mumps, rubella, chickenpox, and influenza. In some locations, some diseases periodically occur every year, such as chickenpox, mumps, and poliomyelitis [1]. As the main reason to investigate that occurrence of this type of bifurcations it that they play a relevant role for disease control and eradication. In fact, there has been more and more interest in the analysis and prediction of consequences of public health strategies designed to control infectious diseases. The asymptotic behavior of epidemic models has been studied by many researchers. But a very different bifurcation behavior has been found in recent papers, see for example, Fred Brauer [2], J. Dushoff et al. [3], Hui and Zhu [4], P. van den Driessche and James Watmough [5], L. Liu, X.G. Li, K.J. Zhuang [6], T.L. Zhang, J.L. Liu, Z.D. Teng [7,8], B.D. Hassard, N.D. Kazarinoffand, Y.H. Wan [9], Carlo B., Massimiliano F., Luca G. [10] and Abid A., Khalid H. [11]. In this paper, from Showroq' studding she studied just local and global stability [12]. In addition to the above studied, we discuss several types of bifurcations: Saddle-node, transcritical, pitchfork, and Hopf bifurcation near all the equilibrium points.

2 Mathematical model

In this section, an epidemic population with a limited resource for treatment involving immigrations is proposed for study. The population is divided into two classes the susceptible individuals $S(t)$ at time t and the infected individuals $I(t)$ at time t . It is assume that a constant flow say A , of new members arrives into the population in unit time with the fraction p ($0 \leq p \leq 1$) of A arriving infected. Accordingly the dynamics of SIS epidemic model with a limited resource for treatment and constant rate of immigrations, which is represented by the following system of nonlinear ordinary equations:

$$\begin{aligned} \dot{S} &= \Lambda + (1-p)A - (\mu + \beta I)S + \gamma I + T(I) \\ \dot{I} &= pA + \beta SI - (\mu + \alpha + \gamma)I - T(I) \end{aligned} \quad (1)$$

Note that, all the parameters $\Lambda, A, \mu, \gamma, \alpha$ and β of system (1) are assumed to be positive and can be described as follows: Λ is the recruitment rate of the susceptible population, A is the constant flow rate of immigrants, μ is the natural death rate in each classes, γ is the nature recovery rate from infected individuals to susceptible individuals, α is the disease related death rate, β is the infected coefficient, finally $T(I)$ is the treatment function which given by [13]:

$$T(I) = \begin{cases} I & \text{if } 0 < I \leq I_0, \\ k & \text{if } I > I_0. \end{cases} \quad (2)$$

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Here, $K = rI_0$ this means that the treatment rate is proportional to the number of infected individuals when the capacity of treatment is not reached, and other wise takes the maximal capacity. Obviously, due to the biological meaning of the components $S(t)$ and $I(t)$ we focus on the model in the domain: $\mathfrak{R}_+^2 = \{(S, I) \in \mathfrak{R}_+^2 : S \geq 0, I \geq 0\}$ which is positively invariant for system (1). Note that, study system (1) when $(0 < I \leq I_0)$ of treatment function only then can be rewritten in the following form:

$$\begin{aligned} \dot{S} &= \Lambda + (1 - p)A - (\mu + \beta I)S + (\gamma + r)I \\ \dot{I} &= pA + \beta SI - (\mu + \alpha + \gamma + r)I \end{aligned} \tag{3}$$

Now, all the solutions of above system which initiate in \mathfrak{R}_+^2 are uniformly bounded and this system has two possible equilibrium points, namely $E_1 = (S_1, 0)$ is called a disease free equilibrium point and the second point is called endemic equilibrium point and denoted by $E_2 = (S_2, I_2)$ that given in [12] where:

$$S_1 = \frac{\Lambda + A}{\mu} \tag{4}$$

$$S_2 = \frac{\Lambda + A(1 - p) + (\gamma + r)I_2}{\mu + \beta I_2} \tag{5}$$

$$I_2 = \frac{-a_2}{2a_1} - \frac{\sqrt{a_2^2 - 4a_1a_3}}{2a_1} \tag{6}$$

here: $a_1 = -\beta(\mu + \alpha)$; $a_2 = \beta(\Lambda + A) - \mu(\mu + \alpha + \gamma + r)$; $a_3 = p\mu A$

3 The local bifurcation analysis

In this section, the effect of varying the parameter on the dynamical behavior of the system (3) around each equilibrium points is studied. Recall that the existence of non-hyperbolic equilibrium point of system (3) is the necessary but not sufficient condition for bifurcation to occur. Therefore, in the following theorems are applications to the Sotomayor's theorem [14] for local bifurcation is adapted.

Theorem (1): The Jacobian matrix of system (3) at (E_1) with parameter:

$$\beta^* = \frac{\mu + \alpha + \gamma + r}{S_1} \tag{7}$$

has:

- 1) No saddle-node bifurcation
- 2) No pitchfork bifurcation
- 3) Transcritical bifurcation

Proof: It is easy to verify that the Jacobian matrix of system (3) at (E_1, β^*) can be written as:

$$J = Df(E_1, \beta^*)$$

$$= \begin{pmatrix} -\mu - \beta^* S_1 + \gamma + r & \\ 0 & 0 \end{pmatrix}$$

Clearly, the second eigenvalue λ_I in the I- direction is zero, while λ_S is negative. Further, the eigenvector (say $Z = (z_1, z_2)^T$) corresponding to λ_I satisfy the following:

$$JZ = \lambda Z$$

then

$$JZ = 0$$

Thus

$$\begin{pmatrix} -\mu - \beta^* S_1 + \gamma + r & \\ 0 & 0 \end{pmatrix} \cdot \begin{pmatrix} z_1 \\ z_2 \end{pmatrix} = 0$$

From which we get that:

$$-\mu z_1 - \beta^* S_1 z_2 + (\gamma + r)z_2 = 0 \tag{8}$$

So by solving the above system of equation we get:

$$z_1 = qz_2$$

where: $q = \frac{-[\beta^* S_1 - (\gamma + r)]z_2}{\mu}$, here z_2 be any non zero real number. Thus

$$Z = \begin{pmatrix} qz_2 \\ z_2 \end{pmatrix}$$

Similarly the eigenvector $W = (w_1, w_2)^T$ corresponding to λ_I of J^T can be written:

$$\begin{pmatrix} -\mu & 0 \\ -\beta^* S_1 + \gamma + r & 0 \end{pmatrix} \cdot \begin{pmatrix} w_1 \\ w_2 \end{pmatrix} = 0$$

This gives:

$$\begin{aligned} -\mu w_1 &= 0 \\ (\gamma + r - \beta^* S_1)w_1 &= 0 \end{aligned} \tag{9}$$

So by solving the above system of equation we get:

$$W = \begin{pmatrix} 0 \\ w_2 \end{pmatrix}$$

here z_2 be any non zero real number. Now rewrite system (3) in a vector form as:

$$\frac{dX}{dt} = f(X)$$

where, $X = (S, I)^T$ and $f = (f_1, f_2)^T$ with $f_i, i = 1, 2$ given in system (3), and then determine $\frac{df}{d\beta} = f_\beta$ we get that:

$$f_\beta = \begin{pmatrix} -SI \\ SI \end{pmatrix}$$

Then

$$f_{\beta}(E_1, \beta^*) = \begin{pmatrix} 0 \\ 0 \end{pmatrix}$$

Therefore:

$$W^T \cdot f_{\beta}(E_1, \beta^*) = 0$$

Consequently, according to Sotomayor's theorem the system has no saddle-node bifurcation near E_1 at β^* . Now in order to investigate the accruing of other types of bifurcation, the derivative of f_{β} with respect to vector X , say $Df_{\beta}(E_1, \beta^*)$, is computed

$$Df_{\beta}(E_1, \beta^*) = \begin{pmatrix} 0 & -S_1 \\ 0 & S_1 \end{pmatrix}$$

So

$$W^T \cdot [Df_{\beta}(E_1, \beta^*) \cdot Z] = w_2 z_2 S_1 \neq 0$$

Again, according to Sotomayor theorem, if in addition to the above the following holds

$$W^T \cdot [D^2 f(E_1, \beta^*) \cdot (Z, Z)] \neq 0$$

Here $Df(E_1, \beta^*)$ is the Jacobian matrix at E_1 and β^* , then the system (3) possesses a transcritical bifurcation but no pitch-fork bifurcation can occur. Now since we have that:

$$[D^2 f(E_1, \beta^*) \cdot (Z, Z)] = \begin{pmatrix} 2q\beta^* z_2^2 \\ -2q\beta^* z_2^2 \end{pmatrix}$$

Therefore:

$$W^T \cdot [D^2 f(E_1, \beta^*) \cdot (Z, Z)] = -2q\beta^* w_2 z_2^2 \neq 0$$

Then the system (3) has a transcritical bifurcation at E_1 when the parameter β passes through the bifurcation value β^* .

Theorem (2): The Jacobian matrix of system (3) at (E_2) with parameter:

$$\alpha_o = \frac{\mu[\beta S_2 - (\mu + \gamma + r + \beta I_2)]}{\mu + \beta I_2} \tag{10}$$

Clearly α_o is positive provided that:

$$\beta S_2 > \mu + \gamma + r + \beta I_2 \tag{11}$$

Then the system (3) has:

- 1) Saddle-node bifurcation
- 2) No pitchfork bifurcation
- 3) No Transcritical bifurcation

Proof: The system (3) at the endemic equilibrium point E_2 has zero eigenvalue (say $\bar{\lambda}$) if and only if $\det(J(E_2)) = 0$, therefore $\alpha_o = \frac{\mu[\beta S_2 - (\mu + \gamma + r + \beta I_2)]}{\mu + \beta I_2}$ is taken as a candidate bifurcation parameter and the Jacobian matrix $J(E_2)$ with $\alpha = \alpha_o$ becomes:

$$J(\bar{\lambda} = 0) = \begin{pmatrix} -(\mu + \beta I_2) & -\beta S_2 + (\gamma + r) \\ \beta I_2 & \beta S_2 - (\mu + \alpha_o + \gamma + r) \end{pmatrix}$$

Now, to find eigenvector (say $L = (l_1, l_2)^T$) corresponding to $\bar{\lambda} = 0$ satisfy the following:

$$J \cdot L = 0$$

Thus:

$$\begin{pmatrix} -(\mu + \beta I_2) & -\beta S_2 + (\gamma + r) \\ \beta I_2 & \beta S_2 - (\mu + \alpha_o + \gamma + r) \end{pmatrix} \cdot \begin{pmatrix} l_1 \\ l_2 \end{pmatrix} = 0$$

From which we get that:

$$-(\mu + \beta I_2)l_1 - (\beta S_2 - (\gamma + r))l_2 = 0 \tag{12}$$

$$\beta I_2 l_1 + (\beta S_2 - (\mu + \alpha_o + \gamma + r))l_2 = 0 \tag{13}$$

So by solving the above system of equations we get:

$$l_1 = z l_2$$

where: $z = \frac{-(\beta S_2 - (\gamma + r))l_2}{\mu + \beta I_2}$ here l_2 be any non zero real number. Thus

$$L = \begin{pmatrix} z l_2 \\ l_2 \end{pmatrix}$$

Similarly the eigenvector $H = (h_1, h_2)^T$ corresponding to $\bar{\lambda} = 0$ of J^T can be written:

$$\begin{pmatrix} -(\mu + \beta I_2) & \beta I_2 \\ -\beta S_2 + \gamma + r & \beta S_2 - (\mu + \alpha_o + \gamma + r) \end{pmatrix} \cdot \begin{pmatrix} h_1 \\ h_2 \end{pmatrix} = 0$$

This gives:

$$-(\mu + \beta I_2)h_1 + \beta I_2 h_2 = 0 \tag{14}$$

$$(\gamma + r - \beta S_2)h_1 + (\beta S_1 - (\mu + \alpha_o + \gamma + r))h_2 = 0 \tag{15}$$

So by solving the above system of equation we get:

$$H = \begin{pmatrix} q h_2 \\ h_2 \end{pmatrix}$$

where: $q = \frac{\beta I_2}{\mu + \beta I_2}$ and h_2 be any non zero real number. Now rewrite system (3) in a vector form as:

$$\frac{dY}{dt} = f(Y)$$

where $Y = (S, I)^T$ and $f = (f_1, f_2)^T$ with $f_i, i = 1, 2$ given in system (3), and then determine $\frac{df}{d\alpha} = f_{\alpha}$ we get that:

$$f_{\alpha} = \begin{pmatrix} 0 \\ -I \end{pmatrix}$$

Then

$$f_{\alpha}(E_2, \alpha_o) = \begin{pmatrix} 0 \\ I_2 \end{pmatrix}$$

Therefore:

$$H^T \cdot f_\alpha(E_2, \alpha_0) = -h_2 I_2 \neq 0$$

So, according to Sotomayor's theorem for local bifurcation the transcritical bifurcation and pitchfork bifurcation cannot occur. While the first condition of the saddle-node bifurcation is satisfied. Moreover it easy to verify that

$$D^2 f(E_2, \alpha_0) \cdot (L, L) = \begin{pmatrix} 2z\beta l_2^2 \\ -2z\beta l_2^2 \end{pmatrix}$$

Hence

$$H^T \cdot [D^2 f(E_2, \alpha_0) \cdot (L, L)] = 2z\beta h_2 l_2^2 (q - 1) \neq 0$$

Consequently, according to Sotomayor's theorem the system (3) has saddle-node bifurcation near E_2 at α_0 .

4 The Hopf bifurcation analysis

In this section, the existence of periodic dynamic in system (3) due to changing the value of one parameter is studied in the following theorem.

Theorem (3): The system (3) has a Hopf-bifurcation around the endemic equilibrium point E_2 satisfy the following condition:

$$\bar{\beta} = \frac{2\mu + \alpha + \gamma + r}{S_2 + I_2} \quad (16)$$

Proof: Consider the Jacobian matrix of system (3) at E_2 with the characteristic equation can be written in the following form:

$$J(E_2) = \begin{pmatrix} -(\mu + \beta I_2) & -\beta S_2 + \gamma + r \\ \beta I_2 & \beta S_2 - (\mu + \alpha + \gamma + r) \end{pmatrix}$$

Then

$$\lambda^2 + T\lambda + D = 0 \quad (17)$$

Clearly, the eigenvalues of above equation can be written:

$$\lambda = \frac{1}{2} \left[-T \pm \sqrt{T^2 - 4D} \right]$$

here

$$T(\text{trace}) = \beta(S_2 - I_2) - (2\mu + \alpha + \gamma + r)$$

$$D(\text{det}) = \mu(\mu + \alpha + \gamma + r - \beta S_2) + \beta I_2(\mu + \alpha + 2(\gamma + r))$$

Obviously, system (3) dose not a Hopf-bifurcation around the endemic equilibrium point E_2 if and only if the trace of eigenvalues $T \neq 0$. Now, the necessary and sufficient conditions for a simple Hopf-bifurcation to occur we need to find a parameter ($\bar{\beta}$) satisfy that:

$$T(\bar{\beta}) = 0 \quad (18)$$

$$\frac{dT}{d\beta} \Big|_{\beta=\bar{\beta}} \neq 0 \quad (19)$$

Then the system (3) has two complex conjugate eigenvalues. Clearly, the first condition (18) for the Hopf-bifurcation is satisfied at $\beta = \bar{\beta}$ if and only if provided the condition (16). Let us now check the second condition (19) in the following:

$$\frac{dT}{d\beta} \Big|_{\beta=\bar{\beta}} = S_2 \neq 0$$

Hence, the system (3) has a Hopf-bifurcation a round the endemic equilibrium point E_2 at the parameter $\beta = \bar{\beta}$ and the proof is complete.

5 Numerical analysis

In this section, the local and Hopf bifurcation of system (3) is studied. The objectives of this study are confirming our analytical results and understand the effects of bifurcation value on the dynamics of SIS epidemic system. Consequently, system (3) is solved numerically for same initial condition (150,550) and different sets of parameters. It is observed that, for the following set of hypothetical parameters, system (3) is solved numerically for same set of initial value and then the trajectories of system (3) as a function of time are drawn in Figure (1).

$$\begin{aligned} \Lambda &= 400; A = 100; p = 0; \beta = 0.0001 \\ \gamma &= 2; \alpha = 2; r = 1; \mu = 0.1 \end{aligned} \quad (20)$$

Obviously, Figure (1) Show clearly the transcritical bifurcation accurse of system (3) near the disease free equilibrium point E_1 , when the infection rate increases (through increasing β), the disease free equilibrium point of system (3) becomes unstable point and the trajectory of system (3) approaches asymptotically to the endemic equilibrium point.

However, for the data given equation (20) but $p = 0.01$ and $\beta = 0.01$. The trajectories of system (3) are drawn in Figure (2).

Clearly, from above figure as the infection rate decrease (β), then the endemic equilibrium point losses it stability and a Hopf bifurcation occurs as shown in Fig. (2a). however, as it increases the endemic equilibrium point still coexists and stable with increase in the value of infected individuals whereas the value susceptible individuals increases.

Now, in order to discuss the effect of varying the treatment rate on the dynamical behavior of system (3), the system is solved for different values of treatment rate $r = 3, 4, 5, 24$ respectively, keeping other parameters fixed as given in equation (20) with $p = 0.01$ and $\beta = 0.01$, and

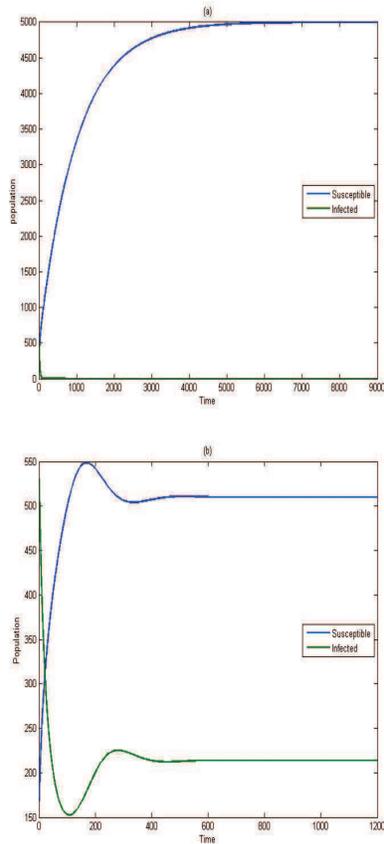


Fig. 1: (a) Time series of trajectories of system (3) for data given in equation (20). (b) Time series of trajectories of system (3) for data given in equation (20) with $\beta = 0.01$.

then the solution of system (3) is drawn in Figure (3).

According to the above figure, it is clear that, as the treatment rate increases the endemic equilibrium point loses its stability and a Hopf bifurcation occurs as shown in Figure (3c). However, as it decreases the endemic equilibrium point still coexists and stable.

6 Conclusions and discussion

In this paper, the system (3) has a transcritical bifurcation near the disease free equilibrium point, but neither saddle-node nor pitchfork bifurcation can accrue. And, has a Hopf bifurcation near the endemic equilibrium point. Finally according to the numerical simulation the following results are obtained:

1) As the varying the infection rate increases, the asymptotic behavior of the system transfers from approaching to disease free equilibrium point to the endemic equilibrium point.

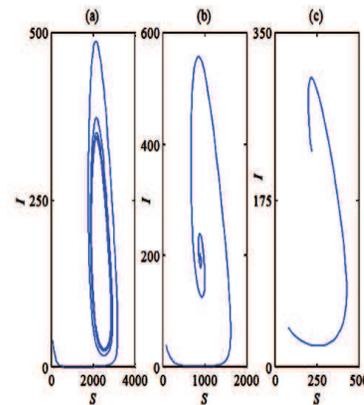


Fig. 2: (a) phase plots of system (3) for data given in equation (20) with $p = 0.01$ and $\beta = 0.002$. (b) Phase plots of system (3) for data given in equation (20) with $p = 0.01$ and $\beta = 0.005$. (c) Phase plots of system (3) for data given in equation (20) with $p = 0.01$ and $\beta = 0.02$.

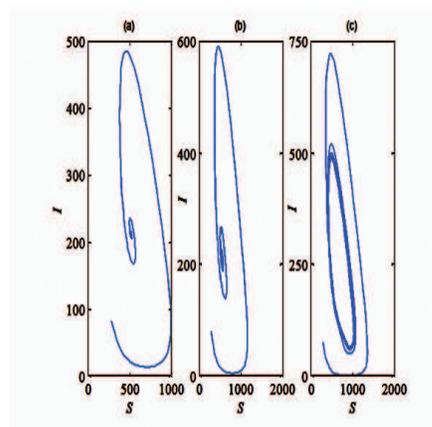


Fig. 3: (a) phase plots of system (3) for data given in equation (20) with $r = 3$. (b) phase plots of system (3) for data given in equation (20) with $r = 4$. (c) phase plots of system (3) for data given in equation (20) with $r = 5.24$.

2) As the infected rate decreases then the system (3) still approaches to the endemic equilibrium point with the system (3) has occurrence of a Hopf bifurcation around the endemic equilibrium point .

3) Finally, the treatment rate increases then the asymptotic behavior of system (3) losses stability and a Hopf bifurcation occurs around the endemic equilibrium point.

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