



## DESIGN AND ANALYSIS OF MODIFIED SIQRS MODEL FOR PERFORMANCE STUDY OF WIRELESS SENSOR NETWORK

RUDRA PRATAP OJHA, GOUTAM SANYAL \*, PRAMOD KUMAR SRIVASTAVA† AND KAVITA SHARMA ‡

**Abstract.** The dynamics of worm propagation in the Wireless Sensor Networks (WSNs) is one of the fundamental challenge due to critical operational constraints. In this paper we propose a modified Susceptible-Infectious-Quarantined-Recovered-Susceptible (SIQRS) model based on epidemic theory. The proposed model demonstrate the effect of quarantined state on worms propagation in WSNs. This model incorporates communication radius, area of communication and the associated node density. The spreading dynamics of worms defined with the help of Basic Reproduction Number ( $R_0$ ) and if  $R_0$  is less than or equal to one the worm-free equilibrium is globally asymptotically stable, and if  $R_0$  is greater than one the worm will persists in the system. This model formulated by differential equations and explain the process of worm propagation in WSNs. We also study the effect of different parameters on the performance of system. Finally, the control mechanism and performance of the proposed model is validated through extensive simulation results.

**Key words:** Epidemic model, Basic reproduction number, Stability, Wireless Sensor Network, Communication radius, Node density

**AMS subject classifications.** 68M10, 90B18

**1. Introduction.** Data communication is one of the primary requirements of modern society. There are different ways of data communication, it may be wired and wireless. The wireless sensor network is also a kind of wireless communication and it having great potential of application like military, patient health monitoring, vehicle traffic monitoring, battlefield, environmental monitoring etc.[1]. Wireless sensor network is a collection of large number of sensor nodes and it is a small device equipped with memory, processing unit, energy source and communication units. The Sensor nodes can be deployed in any type of terrain for collection of data from surroundings and send them to sink node via neighbor nodes. There are some limitations of sensor nodes for example communication range, energy, memory etc. therefore data delivered at the sink node in multi-hop [2]. The Sensor nodes are scattered in a hostile environment with restricted power and charging is very difficult task at deployed location. In inaccessible region, the sensors nodes are deployed by robot or airplane. Due to large applications of WSNs it becomes one of the hot topic for researchers. Energy consumption and increasing the lifetime of WSNs methods has been developed[2, 3, 4], another method related to network topology [5] and placement[6] of nodes has been also studied. WSNs have resource constraints therefore it has a weak defense and soft target for malware attack on nodes [7]. The sensor nodes can be easily targeted by software attacks like virus or worm. Nowadays, wireless devices are targeted by malicious codes very easily and spread from device to device through wireless communication for example Bluetooth and Wi-Fi[8, 9]. The controlling of worm propagation is one of the difficult task. Mathematical modeling is an important tool for analyses the dynamics of worm propagation in wireless sensor network. A worm attack on a certain node in wireless sensor network and its gets infected, then infection spread through the neighbour nodes in the entire network [10]. Therefore, security mechanism against worms attack for wireless sensor network has essential practical importance. Since, there is fundamental similarity between biological worm transmission based on epidemic model and the software generated worm in wireless network.

The social researchers used the concept of epidemic model [11, 12, 13] comprehensively and could be correctly applied to worms spread in WSN. There are some allied applications of epidemic models in the literature of wireless network [8, 13, 14, 15, 16]. The spreading of virus over the internet has been broadly studied by researchers based on the concept of epidemic model [13, 14, 17, 18, 19] and some epidemiological model extensively developed for WSN [8, 10, 16]. A simple algorithm for information diffusion SI model [14] was proposed for analysis of worm propagation in mobile ad hoc networks and developed an expression that describe the relation between rate of infection and node density. Another modified SI model [10] was proposed

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to improve the antivirus capability of networks by leveraging sleep mode and capture both spatial and temporal dynamics simultaneously. This model was verified with the help of extensive simulation.

In this paper, we study the temporal and spatial dynamics and possible attacking behavior of worms in the wireless sensor network. The proposed model depicts worm propagation behavior in wireless sensor network with quarantine state in real world. The meaning of quarantine is to forcefully isolate and stop working of nodes. When infected nodes are detected at any time in the network then they promptly be isolated by detection program. We can stop the spreading of worm and improve the security of network as well as increase the lifetime of wireless sensor network by using this model. Remaining part of the paper is ordered as follows. Section 2 include the related work; Section 3 describes the proposed model. Modeling and analysis of proposed model is given in section 4. Equilibrium and stability studied in section 5. Simulation results and performance analysis in section 6. Section 7 conclusion and future scope.

**2. Related work.** Studies the process of worm propagation on the Internet based on epidemic theory was proposed by [19]. There are some worm propagation conventional model on the internet: SIS, SIR, Two-factor and IWM model [20, 21]. These models comprise of differential equation, and they can successfully portray the qualities of worm spread on the internet. Practically speaking, the SIR model is an expansion of SIS model and it is generally connected in investigating the flow of worm proliferation on the internet. In any case, the above models are not especially intended for WSNs. Therefore, these models do not bolster the worm propagation with the vitality utilization of nodes in WSNs. There are different epidemic models have been examined by various researchers that investigate the dynamic behavior of worm spreading and controlling in WSNs.

In [22] the author study the virus spreading behavior of SIRS model in the wireless sensor network that includes the degree of nodes, topological connection and develops the method for the prevention and controlling of local infection.

Pietro et al. [23] proposed an epidemic model for information survivability in unattended WSNs to ensure the quality of service and node energy in presence of attacker.

In [24] presented a SI model with inclusion of communication radius and node density that illustrate the effect of these parameters on spreading of virus in wireless sensor network with MAC mechanism. But this model did not include the recovery of nodes and other method of protection due to attack of virus, overcome this problem in [25] introduced a model that was SIR-M in which one more state recovery included, and carry out maintenance task in the sleep state of node and recover the infected node without any overhead and enhance the lifetime of WSNs and model was demonstrated by the simulation.

Author [26] proposed an improved SIRS model with inclusion of dead node that describe the process of worm propagation and also included the energy consumption of nodes in WSNs. But this model did not include a number of significant characteristics of model for example reproduction number, stability condition, equilibrium points and delay of network. These missing uniqueness are discussed by some different authors. In [27] author calculate the reproduction number and examines its effect on the wireless sensor network as well as also found the condition of worm free equilibrium and local stability condition of the proposed model. This model was verified by simulation with the help of MATLAB. Zhu and Zhao [28] have discussed another model with inclusion of dead node to study the behavior of malware propagation in wireless sensor network. Numerical proof and simulation shows the dynamic behavior of worm propagation in wireless sensor network but this model do not focus on characteristics and specifications of particular sensor node. These issues are talked by [29], proposed an individual-based models that determine specific attributes of each component of model and beat the deficiency of the past model by the consideration of each device characteristics and evaluate each individually. Ojha et al. [30] proposed a model to study the effect of undetected infectious nodes and effect of treatment. They also, calculate the threshold value and analyze its effect on the wireless sensor network and discuss the local and global equilibrium stability of worm free state. These are proved with the help of simulation.

In [31] a control mechanism developed to efficiently hold down the worm outbreak and escape the network breakdown. This model includes number of nodes, transmission range of sensors, node density etc. and verified by simulation. Keshri and Mishra [32] proposed an epidemic model with inclusion of the two delay period and study its effect on different class of the model. They also found the local and global stability of worm free equilibrium point of the system and calculate its reproduction number of the proposed system as well as observed its effect on wireless sensor network, communication radius, node density and quarantined class have not

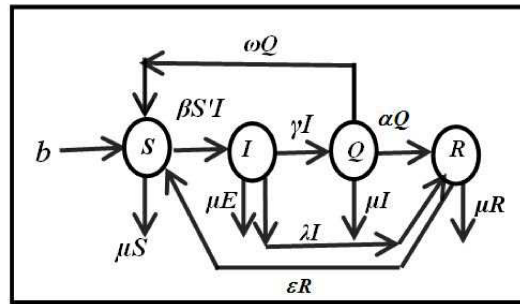


FIG. 3.1. Transition relationship of node states

TABLE 3.1  
Parameters Used

S.No	Symbol	Meaning of Symbol
1	$S$	Susceptible
2	$I$	Infected
3	$Q$	Quarantined
4	$R$	Recovered
5	$b$	Birth rate
6	$\mu$	Death rate
7	$\beta$	Rate of infection
8	$\alpha$	Probability of moving from exposed node to infectious category
9	$\gamma$	Recovery rate of an infectious node
10	$\omega$	Probability of moving from Quarantined node to Susceptible node
11	$\epsilon$	probability at which a recovered node become a susceptible node

been considered in proposed model. A SIRS model was proposed by [33] in which consider the communication radius and node density as well as calculate the reproductive number and found the Eigen values for stability verification of the system. But this model have not efficient mechanism to control the quick spreading of worm in the network. We extend the proposed model with introduction of quarantined class. The quarantined state plays a significant role and sends the infectious nodes into isolation mode and removes the worm from network without any overhead and stretch the life span of network. This model overcomes the insufficiency of existing models.

**3. The proposed model.** We consider the total numbers of nodes in the network are  $N(t)$  at any time  $t$  which is uniformly distributed in the region  $L \times L$  with average density  $\rho$  and radio range of sensor node is  $r$ . There are four states in the proposed model-Susceptible state  $S$  -The nodes which are not infected but vulnerable to the worm. Infected state  $I$  -The infected node which is capable to infecting other nodes. The Quarantined  $Q$  -The isolation state of node and Recovered state  $R$  -The recovered nodes become immune. In the proposed model, we have assumed that the deployment of sensor nodes are uniform and random in two dimensional areas. Each sensor nodes having a radio range  $r$  and coverage area of each node is  $\pi \times r^2$ . The significance of coverage area is to at least one sensor node lies within the sensing range because data delivered to the sink through these nodes. Each node detects the events and reported to the sink. Initially, consider each and every node to be susceptible and can be attack by worm. The relationship between the transitions states is depicted by Fig. 3.1. There is some limitation of the model because all sensor nodes are homogeneous.

**4. Modeling and Analysis.** We assume that the communication area of a sensor node is  $\pi r^2$  and the density of a susceptible node in a unit area is  $\rho(t) = S(t)/(L \times L)$ . The total number of neighboring nodes which lie in the communication range of a sensor node is given as :  $S'(t) = \rho(t)\pi r^2 = \frac{S(t)\pi r^2}{L \times L}$ . According to Fig 3.1, we consider following mathematical model of worm propagation.

$$\left. \begin{aligned} \frac{dS}{dt} &= b - \frac{\pi r^2}{L^2} \beta SI + \omega Q + \varepsilon R - \mu S \\ \frac{dI}{dt} &= \frac{\pi r^2}{L^2} \beta SI - (\mu + \lambda + \gamma) I \\ \frac{dQ}{dt} &= \gamma I - (\alpha + \omega + \mu) Q \\ \frac{dR}{dt} &= \alpha Q + \lambda I - (\mu + \varepsilon) R \end{aligned} \right\} \tag{4.1}$$

Let  $\zeta = \frac{\pi r^2}{L^2} \beta$ . Then the above system (4.1) can be written as:

$$\left. \begin{aligned} \frac{dS}{dt} &= b - \zeta SI + \omega Q + \varepsilon R - \mu S \\ \frac{dI}{dt} &= \zeta SI - (\mu + \lambda + \gamma) I \\ \frac{dQ}{dt} &= \gamma I - (\alpha + \omega + \mu) Q \\ \frac{dR}{dt} &= \alpha Q + \lambda I - (\mu + \varepsilon) R \end{aligned} \right\} \tag{4.2}$$

**5. Existence of positive equilibrium.** For equilibrium points we have

$$\frac{dS}{dt} = 0, \frac{dI}{dt} = 0, \frac{dQ}{dt} = 0, \frac{dR}{dt} = 0$$

and after a straight forward calculation, we get equilibrium points as  $P_0 = (S_0, I_0, Q_0, R_0) = (\frac{b}{N}, 0, 0, 0)$  for worm free state and  $P^* = (S^*, I^*, Q^*, R^*)$  for endemic state, where

$$S^* = \frac{\lambda + \gamma + \mu}{\zeta} I^* = (1 - \frac{1}{R_0}) \frac{b(\varepsilon + \mu)(\mu + \alpha + \omega)}{A}, Q^* = (1 - \frac{1}{R_0}) \frac{\gamma(\varepsilon + \mu)}{A}, R^* = (1 - \frac{1}{R_0}) \frac{\alpha(\gamma + \lambda) + \lambda(\mu + \omega)}{A}$$

where  $A = (\mu + \alpha)(\lambda + \mu) + (\mu + \varepsilon) + (\gamma + \varepsilon + \omega) + (\alpha\gamma + \omega\lambda)$  and  $R_0$  [34] is the basic reproduction number and given by  $R_0 = \frac{b\zeta}{\mu(\mu + \lambda + \gamma)}$ . It is clear that  $P^*$  exist and unique if and only if  $R_0 > 1$ .

**5.1. Stability analysis of the worm free equilibrium.**

**THEOREM 5.1.** *The worm free equilibrium (W F E) is locally asymptotically stable if  $R_0 < 1$  otherwise unstable for  $R_0 > 1$ .*

*Proof.* In order to check the stability of point ,we will find all the Eigenvalue values of the variational matrix is given by

$$J(P_0) = \begin{bmatrix} -\zeta I_0 - \mu & \zeta S_0 & \omega & -\varepsilon \\ \zeta I_0 & \zeta S_0 - (\lambda + \gamma + \mu) & 0 & 0 \\ 0 & \gamma & -(\alpha + \mu + \omega) & 0 \\ 0 & 0 & \alpha & -(\varepsilon + \mu) \end{bmatrix} \tag{5.1}$$

□

Eigen values of (5.1) are:  $\theta_1 = -\mu$ ,  $\theta_2 = -(\mu + \alpha + \omega)$ ,  $\theta_3 = (\varepsilon + \mu)$ ,  $\theta_4 = (R_0 - 1)(\frac{1}{\gamma + \lambda + \mu})$ .

It is clear that all eigen values of the variational matrix are negative when  $R_0 < 1$ . Therefore, the system is locally asymptotically stable at worm free equilibrium. Furthermore, the following theorem holds.

**THEOREM 5.2.** *The worm free equilibrium is said to be globally asymptotically stable if  $R_0 \leq 1$ .*

*Proof.* Consider the Lyapunov function  $L(t): R^4 \rightarrow R^+$  defined by  $L(t) = \eta I(t)$ , now taking time derivative of  $L(t)$  with respect to time  $t$  is  $\frac{dL(t)}{dt} = \eta I(t) \Rightarrow \frac{dL(t)}{dt} = \eta(\zeta SI) - (\gamma + \mu + \lambda)I = \eta(\zeta S - (\gamma + \mu + \lambda))I$ . After suitable assumption of  $\eta = \frac{1}{(\gamma + \lambda + \mu)}$  we get  $\frac{dL}{dt} = [\zeta S - (\gamma + \lambda + \mu)] = [\frac{\zeta b}{(\mu + \gamma + \lambda)\mu} - 1]I = (R_0 - 1)I$ . It is clear that  $\frac{dL}{dt} = 0$  only when  $I = 0$ . Therefore, the maximum invariant set  $\Gamma = (S, I, Q, R) \in R_4^+$  is the singleton set. Therefore, the global stability of worm free equilibrium  $P_0$  when  $R_0 \leq 1$  follows from Lasalle invariance principle [35].  $\square$

**5.2. Stability analysis of Endemic equilibrium.**

**THEOREM 5.3.** *Worm endemic state (WES) is locally asymptotically stable if  $R_0 > 1$ .*

*Proof.* In order to check the stability at point  $P^*$  we will find all the Eigen values of the variational matrix:

$$J(P^*) = \begin{bmatrix} -\zeta I^* - \mu & \zeta S^* & \omega & \varepsilon \\ \zeta I^* & \zeta S^* - (\lambda + \gamma + \mu) & 0 & 0 \\ 0 & \gamma & -(\alpha + \mu + \gamma) & 0 \\ \omega & 0 & 0 & -(\varepsilon + \mu) \end{bmatrix} \tag{5.2}$$

Therefore, the corresponding characteristic equation for the above matrix is given by

$$\nu^4 + D_1\nu^3 + D_2\nu^2 + D_3\nu + D_4 = 0 \tag{5.3}$$

where

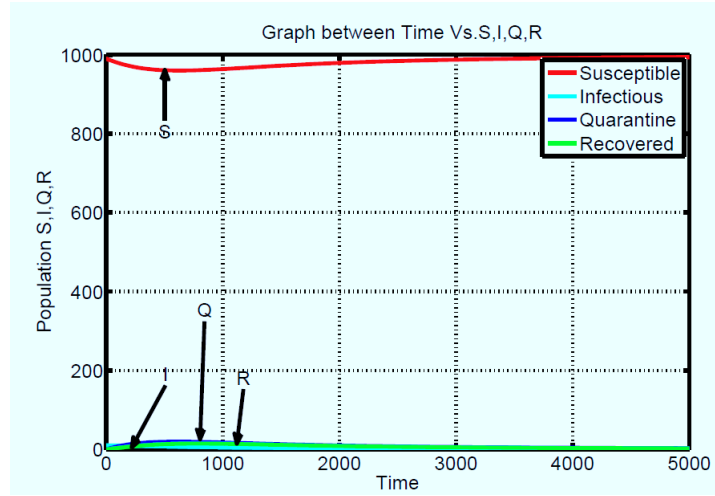
$$\begin{aligned} D_1 &= \zeta Q_1(R_0 - 1) + 3\mu + \alpha + \omega + \varepsilon \\ D_2 &= (\mu + \alpha + \omega)(\mu + \varepsilon) + (\zeta Q_1((R_0 - 1) + \mu)(\alpha + 2\mu + \omega + \varepsilon) + Q_2 \\ D_3 &= \zeta Q_1(R_0 - 1)(\omega\gamma + \varepsilon\lambda) + (\mu + \alpha + \omega)(\mu + \varepsilon)(Q_2 - 1) \\ D_4 &= \zeta Q_1(R_0 - 1)(\gamma\varepsilon\alpha + \omega\gamma(\mu + \varepsilon) + Q_3) \end{aligned}$$

where

$$\begin{aligned} Q_1 &= \frac{b(\mu + \alpha + \omega)(\mu + \varepsilon)}{AR_0}, \\ Q_2 &= -\zeta^2 I^* S^*, \\ Q_3 &= -(\alpha + \mu + \omega)[(\mu + \gamma)(\mu + \varepsilon) + \mu\lambda] \end{aligned}$$

It is clear that all  $D_1, D_2, D_3, D_4$  are positive when  $R_0 > 1$ , by a direct calculation  $G_1 = D_1 > 0, G_2 = D_1 D_2 - D_3 > 0, G_3 = D_3 G_2 - D_1^2 D_4 > 0$  and  $G_4 = D_4, G_3 > 0$ . Therefore, according to Routh Hurwitz criteria, all the roots of equation (5.3) have negative real parts. Therefore, the endemic equilibrium  $P^*$  is locally asymptotically stable when  $R_0 > 1$ . This completes the proof.  $\square$

**6. Simulation and Performance analysis.** It has been shown in the above analysis that  $R_0$  is a threshold value. When  $R_0$  less than equal to one, the fraction of infected nodes vanishes and when  $R_0 > 1$ , the fraction of infected nodes persist in the system. We analyzed the behavior of worm propagation to study the effect of parameters on performance. The proposed model has been simulated with the help of MATLAB and obtained results analyzed below.



$N = 1000; b = 1; L = 10; \beta = 0.0002; \mu = 0.001; \gamma = 0.002; \varepsilon = 0.0003; \omega = 0.0001; \alpha = 0.002; \lambda = 0.0005; r = 0.7$

FIG. 6.1. *Dynamic behavior of model when communication radius is 0.7*

**6.1. Communication Radius of nodes  $r$ .** As we have shown that

$$R_0 = \frac{b\zeta}{\mu(\mu + \gamma + \lambda)}, \text{ where } \zeta = \frac{\pi r^2 \beta}{L^2}.$$

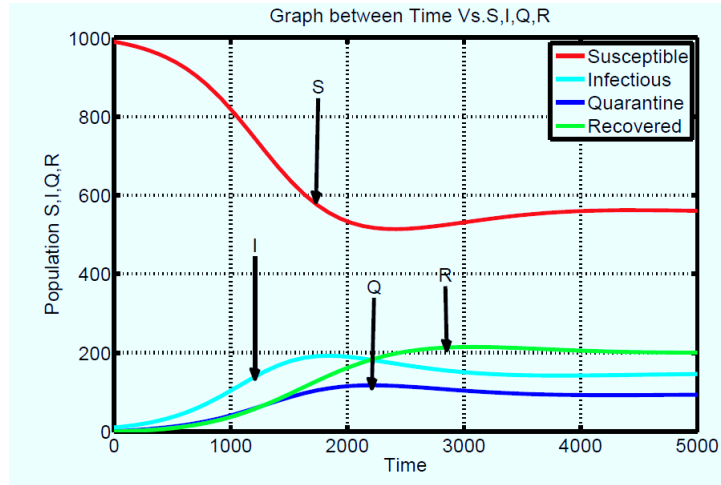
From this equation, we find threshold radius  $r_c = L\sqrt{\frac{(\mu + \gamma + \lambda)\mu}{\pi\beta}}$  i.e when  $r \leq r_c, R_0 \leq 1$  from theorem 5.2, it is clear that the worms in wireless sensor networks can be eradicated and system will stable at worm free equilibrium, when  $r > r_c, R_0 > 1$  according to theorem 5.3, worms in wireless sensor networks will be present consistently and system will stabilize at the endemic equilibrium.

We take following values of parameters as  $N = 1000; L = 10; b = 1; \beta = 0.0002; \mu = 0.001; \gamma = 0.002; \varepsilon = 0.0003; \omega = 0.0001; \alpha = 0.002; \lambda = 0.0005$ . After calculation, we get  $r_c = 0.746542$  the initial values of susceptible, exposed ,infected and recovered nodes in wireless sensor networks are  $S(0) = 990, I(0) = 10, Q(0) = 0$  and  $R(0) = 0$  and takes different values of  $r$ , simulation results are described in Fig.6.1, 6.2 and 6.3. When  $r = 0.7 < r_c$ , Fig.6.1 shows that system (4.2) stabilizes at worm free equilibrium and the simulation results are consistent with theorem 5.2.

When  $r = (1.0 \text{ and } 1.4) > r_c$ , Figs. 6.2 and 6.3 shows that system (4.2) stabilizes at the endemic equilibrium and the simulation results are consistent with theorem 5.3. Communication radius plays an important role for connectivity. Therefore, connectivity is a function of communication radius. From Figs. 6.1, 6.2 and 6.3 we see that when communication radius of nodes increases connectivity increases but the value of  $R_0$  also increases. When the communication radius less than  $r_c$  the network will be stable and worm free. As the communication radius increases  $R_0$  also increases and lead to the failure of the system.

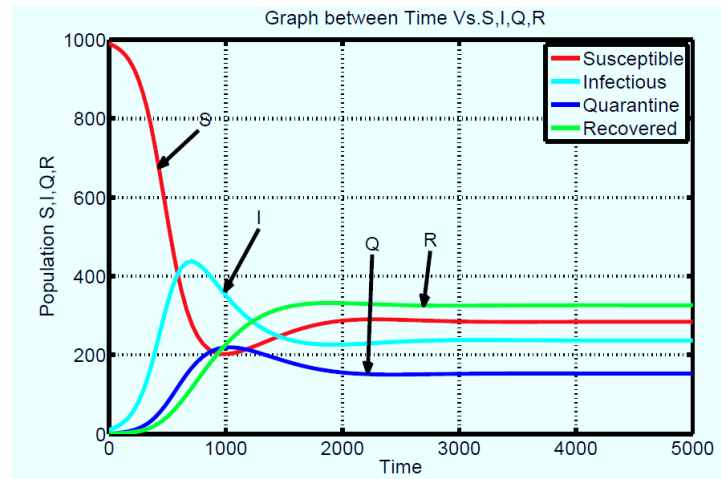
Fig. 6.4 shows that relation between infected and recovered nodes with variation in the parameters. We observed that when the value of  $\alpha$  increases the more number of nodes are recovered smoothly. We also find that if we increase the value of  $\gamma$ , less number of nodes get infected and recovered in short time. This experiment shows the effect of quarantined state on sensor network.

Fig. 6.5 shows the response of the number of susceptible nodes with respect to change in the parameters. We observed that gradually the number of susceptible nodes decrees and recovered nodes increases.



$N = 1000; b = 1; L = 10; \beta = 0.0002; \mu = 0.001; \gamma = 0.002; \varepsilon = 0.0003; \omega = 0.0001; \alpha = 0.002; \lambda = 0.0005; r = 1.0$

FIG. 6.2. Dynamic behavior of model when communication radius is 1.0



$N = 1000; b = 1; L = 10; \beta = 0.0002; \mu = 0.001; \gamma = 0.002; \varepsilon = 0.0003; \omega = 0.0001; \alpha = 0.002; \lambda = 0.0005; r = 1.4$

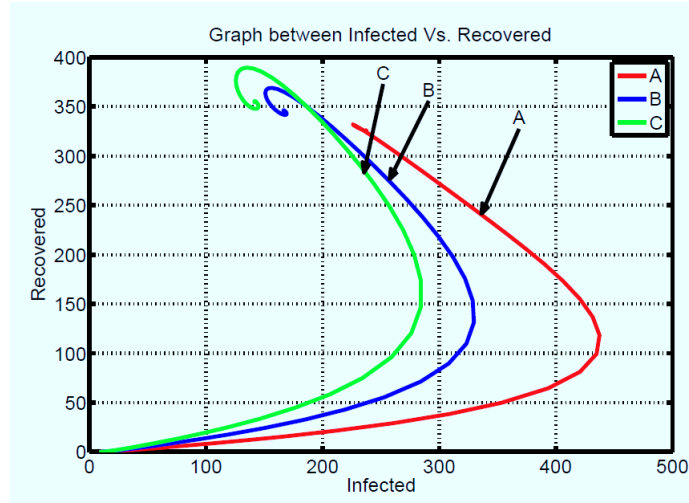
FIG. 6.3. Dynamic behavior of model when communication radius is 1.4

**6.2. Node distributed density.** The threshold value of node density  $\rho_{th}$  is given by

$$\rho_{th} = \frac{N(\mu + \gamma + \lambda)\mu}{b\pi r^2 \beta}$$

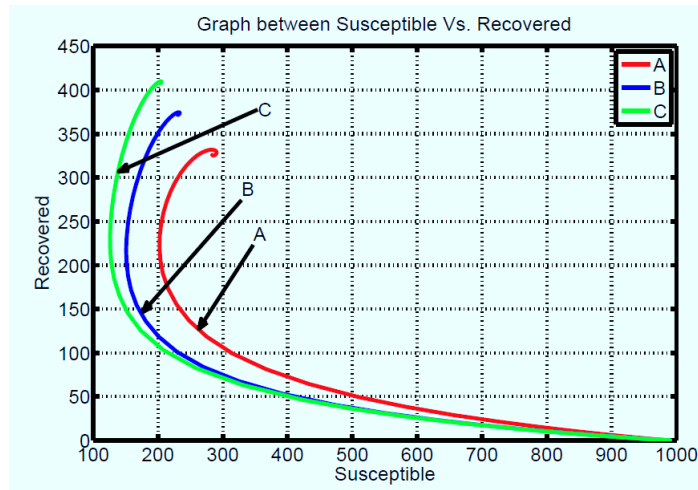
i.e., when  $\rho \leq \rho_{th}, R_0 \leq 1$  then the system has only equilibrium and is globally asymptotically stable; when  $\rho > \rho_{th}, R_0 > 1$ , system has only one endemic equilibrium which is locally asymptotically stable.

We take following values of parameter-  $N = 1000; b = 1; \beta = 0.0002; \mu = 0.001; \gamma = 0.002; \varepsilon = 0.0003; \omega = 0.0001; \alpha = 0.002; \lambda = 0.0005; r = 1$ . After calculation, we get  $\rho_{th} = 5.573248$ . Initial values of susceptible, exposed, infected and recovered nodes in wireless sensor networks are  $S(0) = 990, I(0) = 10, Q(0) = 0$  and  $R(0) = 0$  and When  $L = 11.18$  and  $8.45$ , we can get  $\rho = 8$  and  $14$  using these various values, simulation results are shown in Figs. 6.6 and 6.7. When  $\rho < \rho_{th}$ , the system (4.2) stabilizes at worm free equilibrium and the simulation results are consistent with the theoretical analysis. Whereas when  $\rho = (8 \text{ and } 14) > \rho_{th}$ , Fig. 6.6



$N = 1000; b = 1; L = 10; \beta = 0.0002; \mu = 0.001; \gamma = 0.002; \varepsilon = 0.0003; \omega = 0.0001; \alpha = 0.002; \lambda = 0.0005; r = 1.4$   
 $N = 1000; b = 1; L = 10; \beta = 0.0002; \mu = 0.001; \gamma = 0.003; \varepsilon = 0.0003; \omega = 0.0001; \alpha = 0.003; \lambda = 0.0005; r = 1.4$   
 $N = 1000; b = 1; L = 10; \beta = 0.0002; \mu = 0.001; \gamma = 0.0035; \varepsilon = 0.0003; \omega = 0.0001; \alpha = 0.004; \lambda = 0.0005; r = 1.4$

FIG. 6.4. Plot between Infected and Recovered



$N = 1000; b = 1; L = 10; \beta = 0.0002; \mu = 0.001; \gamma = 0.002; \varepsilon = 0.0003; \omega = 0.0001; \alpha = 0.002; \lambda = 0.0005; r = 1.4$   
 $N = 1000; b = 1; L = 10; \beta = 0.00024; \mu = 0.001; \gamma = 0.003; \varepsilon = 0.0002; \omega = 0.00014; \alpha = 0.0024; \lambda = 0.0004; r = 1.4$   
 $N = 1000; b = 1; L = 10; \beta = 0.00028; \mu = 0.001; \gamma = 0.0035; \varepsilon = 0.00025; \omega = 0.00018; \alpha = 0.003; \lambda = 0.0005; r = 1.4$

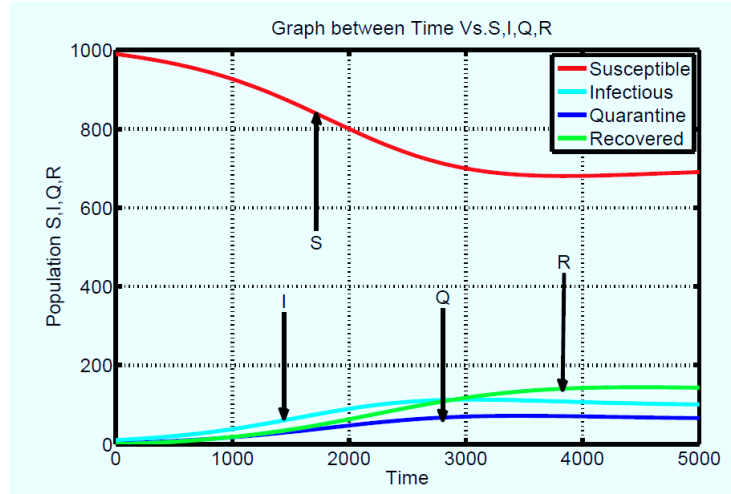
FIG. 6.5. Plot between Susceptible and Recovered

and 6.7 again shows that the trajectories converge to the endemic equilibrium and the simulation results are consistent with the theoretical analysis.

Node density also affects the connectivity of wireless sensor network. From Figs. 6.6 and 6.7 when node density increases connectivity increases and value of  $R_0$  also increases. When the node density less than  $\rho_{th} = 7.043116$  worm do not persist in the network. When the node density increases  $R_0$  also increases and this is harmful for network.

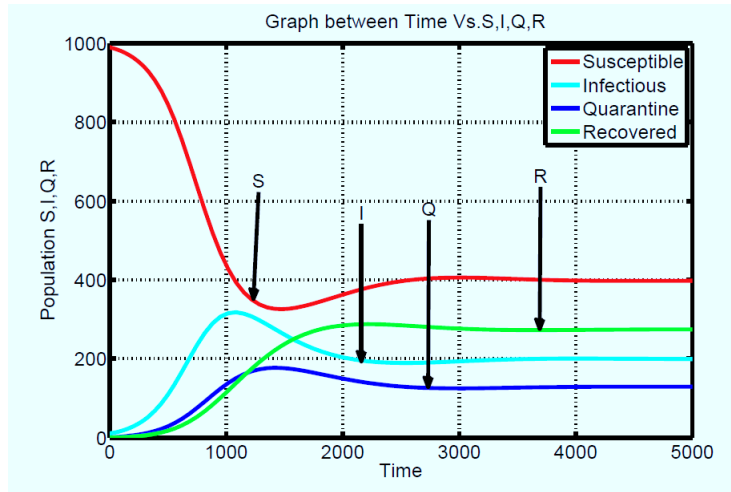
**6.3. Performance analysis of the model.** Figs. 6.10, 6.11 and 6.12 shows the effect of parameters  $\beta$ ,  $\alpha$  and  $\lambda$  on the number of infectious nodes with change of values. As anticipated, when the value of  $\beta$





$N = 1000; \rho = 8; b = 1; \beta = 0.0002; \mu = 0.001; \gamma = 0.002; \varepsilon = 0.0003; \omega = 0.0001; \alpha = 0.002; \lambda = 0.0005; r = 1$

FIG. 6.6. Dynamic behavior of model when node density is 8.0

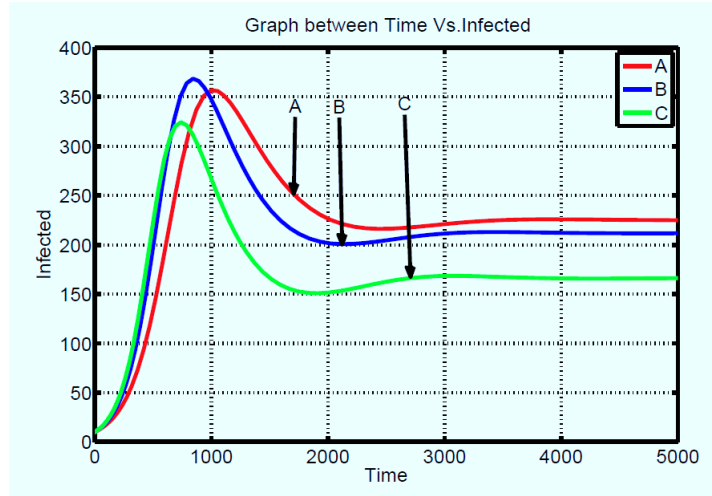


$N = 1000; \rho = 14; b = 1; \beta = 0.0002; \mu = 0.001; \gamma = 0.002; \varepsilon = 0.0003; \omega = 0.0001; \alpha = 0.002; \lambda = 0.0005; r = 1$

FIG. 6.7. Dynamic behavior of model when node density is 14.0

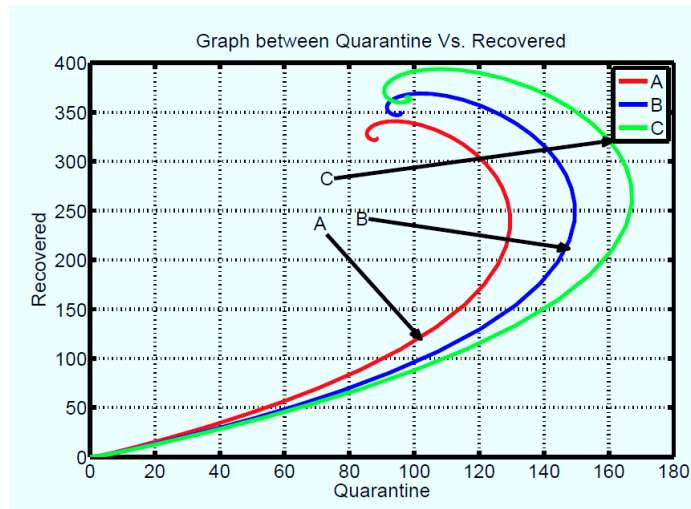
increased, the number of infectious nodes also increased. When  $\lambda$  increased and the remaining parameters are fixed number of recovered nodes increased. It is also found that, as time passes the number of infectious nodes increases and reached at maximum value at a certain stage this will depend on the parameters. Fig. 6.12 shows the comparison between [33] and proposed model. The plot shows that the proposed model is better in comparison to the existing model because more number of nodes are infected in existing model.

**7. Conclusion.** In this paper, we proposed an SIQRS model which describe the spread of worms in wireless sensor network. A controlling parameter  $R_0$  is obtained, which played an important role to understand the dynamic behavior of worms propagation. The worms propagation can be predicted and eradicated from the system that depends on the value of  $R_0$ . We showed that the worm free and equilibrium is locally and globally asymptotically stable, when the value of reproduction number is less than or equal to one and endemic equilibrium is locally asymptotically stable when value of reproduction number is greater than one. Calculate the threshold value of communication radius and node density as well as establish the relation with reproduction



$N = 1000; \beta = 0.0003; \mu = 0.001; \gamma = 0.002; \varepsilon = 0.0003; \omega = 0.0001; \alpha = 0.004; \lambda = 0.0003; r = 1.4; \rho = 4.917$   
 $N = 1000; \beta = 0.00035; \mu = 0.001; \gamma = 0.0023; \varepsilon = 0.0003; \omega = 0.00014; \alpha = 0.0045; \lambda = 0.00034; r = 1.4; \rho = 4.917$   
 $N = 1000; \beta = 0.0004; \mu = 0.001; \gamma = 0.003; \varepsilon = 0.0003; \omega = 0.00018; \alpha = 0.005; \lambda = 0.0005; r = 1.4; \rho = 4.917$

FIG. 6.8. Plot between Time and Infected nodes when node density is 8



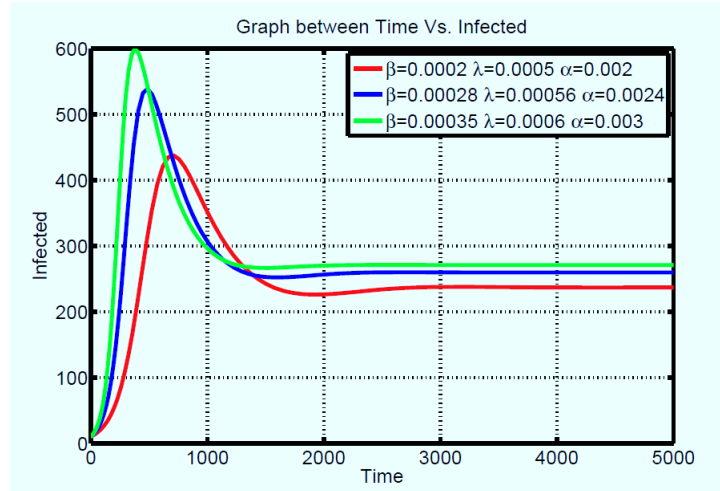
$N = 1000; \beta = 0.0003; \mu = 0.001; \gamma = 0.002; \varepsilon = 0.0003; \omega = 0.0001; \alpha = 0.004; \lambda = 0.0003; r = 1.4; \rho = 4.917$   
 $N = 1000; \beta = 0.0003; \mu = 0.001; \gamma = 0.002; \varepsilon = 0.0003; \omega = 0.00014; \alpha = 0.0045; \lambda = 0.0003; r = 1.4; \rho = 4.917$   
 $N = 1000; \beta = 0.0003; \mu = 0.001; \gamma = 0.003; \varepsilon = 0.0003; \omega = 0.00018; \alpha = 0.005; \lambda = 0.0003; r = 1.4; \rho = 4.917$

FIG. 6.9. Plot between Time and Quarantine nodes when node density is 4.917

number. Furthermore, we have done the simulation of proposed model with the help of MATLAB to verify and validate the results of proposed model. We study the impact of various parameters under different conditions. This model helps to developing an anti-virus mechanism for WSNs. Heterogeneous and moving nodes can be included for analysis of the model in future. On the basis of analysis worm riddance technique can be suggested.

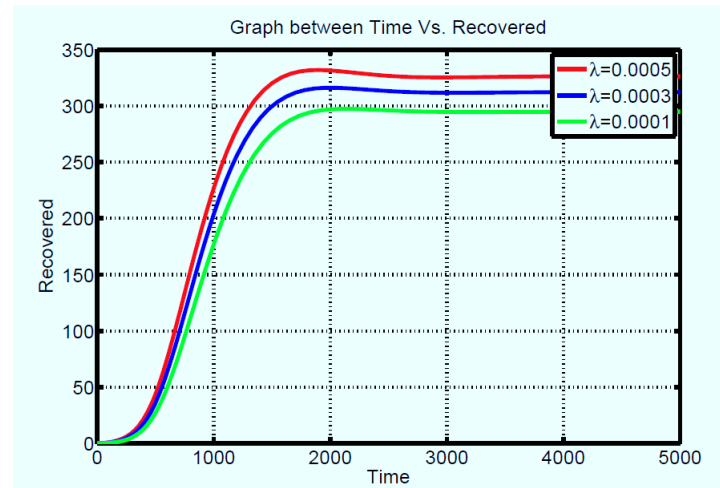
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$N = 1000; b = 1; L = 10; \beta = 0.0002; \mu = 0.001; \gamma = 0.002; \varepsilon = 0.0003; \omega = 0.0001; \alpha = 0.002; \lambda = 0.0005; r = 1.4$   
 $N = 1000; b = 1; L = 10; \beta = 0.00028; \mu = 0.001; \gamma = 0.002; \varepsilon = 0.0003; \omega = 0.0001; \alpha = 0.0024; \lambda = 0.00056; r = 1.4$   
 $N = 1000; b = 1; L = 10; \beta = 0.00035; \mu = 0.001; \gamma = 0.002; \varepsilon = 0.0003; \omega = 0.0001; \alpha = 0.003; \lambda = 0.0006; r = 1.4$

FIG. 6.10. Plot between Time and Infected nodes with variation of parameters



$N = 1000; b = 1; L = 10; \beta = 0.0002; \mu = 0.001; \gamma = 0.002; \varepsilon = 0.0003; \omega = 0.0001; \alpha = 0.002; \lambda = 0.0005; r = 1.4$   
 $N = 1000; b = 1; L = 10; \beta = 0.0002; \mu = 0.001; \gamma = 0.002; \varepsilon = 0.0003; \omega = 0.0001; \alpha = 0.002; \lambda = 0.0003; r = 1.4$   
 $N = 1000; b = 1; L = 10; \beta = 0.0002; \mu = 0.001; \gamma = 0.002; \varepsilon = 0.0003; \omega = 0.0001; \alpha = 0.002; \lambda = 0.0001; r = 1.4$

FIG. 6.11. Plot between Time and Recovered nodes with variation of parameters

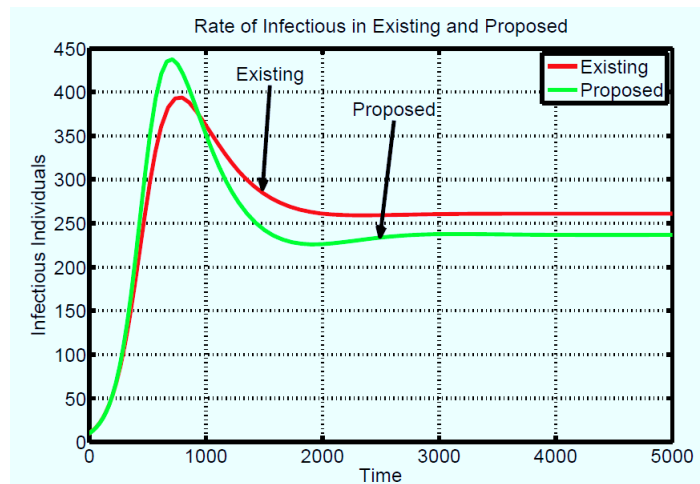
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$N = 1000; b = 1; L = 10; \beta = 0.0002; \mu = 0.001; \gamma = 0.002; \varepsilon = 0.0003; \omega = 0.0001; \alpha = 0.002; \lambda = 0.0005; r = 1.4$

FIG. 6.12. Plot between Time Vs. Infected nodes For Existing and Proposed model

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