Delay Epidemic Model with and without Vaccine

B. Kaymakamzade¹, E. Hincal²

¹ Department of Mathematics, Near East University, Nicosia, TRNC, Mersin 10, Turkey bilgen.kaymakamzade@neu.edu.tr

² Department of Mathematics, Near East University, Nicosia, TRNC, Mersin 10, Turkey evren.hincal@neu.edu.tr

Abstract: In this work, two models with and without vaccine are constucted. For both model basic reproduction numbers are founded as a

$$R_0^1 = \frac{e^{-\mu\tau}\beta\Lambda}{\gamma + \mu + d}, R_0^2 = \frac{e^{-\mu\tau}\Lambda}{(\gamma + \mu + d)(\mu + r)} \left(\beta + \frac{kr}{\mu + 1 - k}\right)$$

From the basic reproduction ratios it can be seen that when there is no vacccine, disease can only controlled with reducing the infectious rate β or decreasing the incubation period. However, when there is a vaccine it is enough to increase the rate of the vaccine r to controlled the disease. In addition to this if transmission rate of infectious also decrease then the disease will disappear faster.

Delay effect on this model is very rare. When we are talking about delay on this paper we mean that incubation period. If we have enough vaccine the effect of delay is very tiny. However if there is no vaccine you can see the effect of delay.

Two equilibria which are disease free and endemic equilibriums are found and with using Lyapunov function it is shown that the global stabilities of each equilibria for both model. For the first model it is found that DFE E_0 is globally asymptotically stable when $R_0^1 < 1$ and endemic equilibrium E_1 is always asymptotically stable. With using similar method it is shown that E_0 is asymptotically stable when $R_0^2 < 1$ and E_1 is always global asymptotically stable for model 2. In last section numerical simulations are given for both model.

Keywords: Delay, Epidemic, Modelling, Vaccine differential equation.

2010 Mathematics Subject Classification: 35N05, 37C75, 34K50

References

[1]

- [2] S.Chauhan, O.P. Misra, J.Dhar. American Journal of Computational and Applied Mathematics 2014, 17-23
- [3] B.Kaymakamzade, E.Hincal. Quality and Quantity DOI 10.1007/s11135-017-0647-8122017
- [4] M.E. Alexander, C. Bowman, S.M. Moghadas, R. Summers, A.B. Gumel, B.M. SahaiA vaccination model for transmission dynamics of influenza SIAM J. Appl. Dyn. Syst., 3 (2004), pp. 503-524
- [5] B. Buonomo, A. dOnofrio, D. LacitignolaGlobal stability of an SIR epidemic model with information dependent vaccination Math. Biosci., 216 (2008), pp. 9-16