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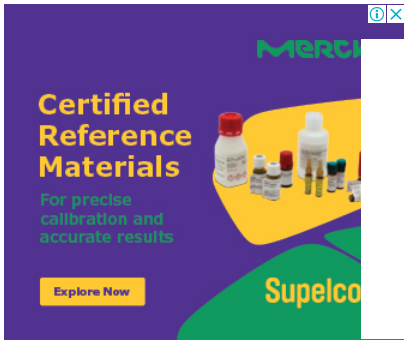
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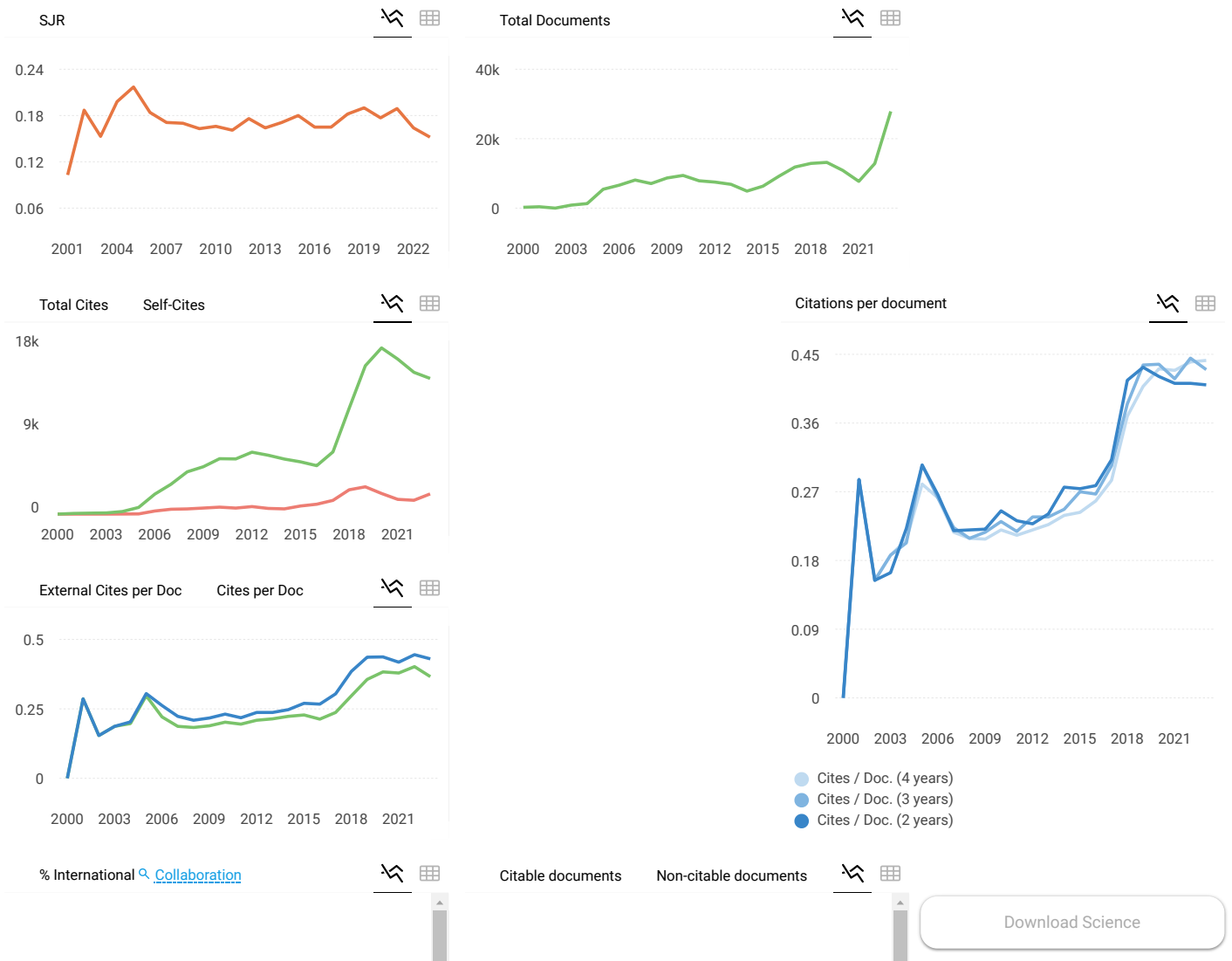
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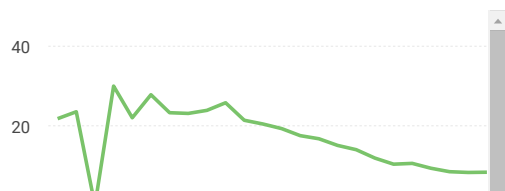


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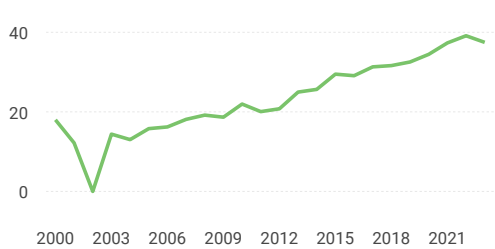
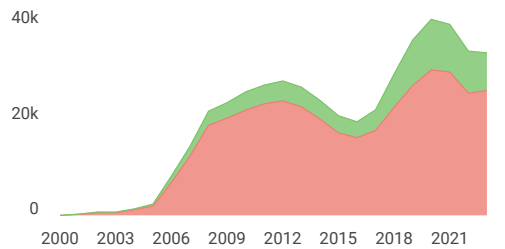




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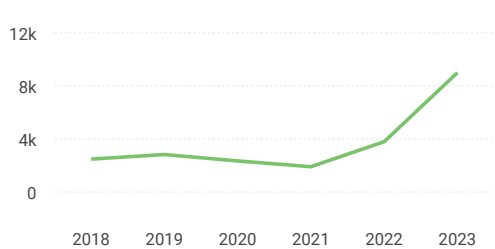
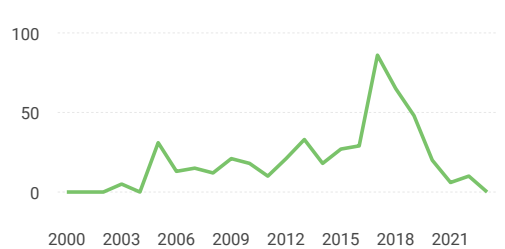
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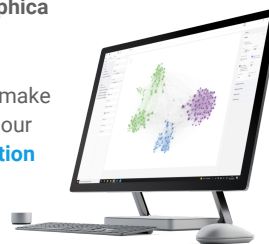
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Yazeed Alzahrani 4 months ago

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The 7th International Conference on Science and Applied Science (ICSAS 2022)

Surakarta, Indonesia • 21 October 2022

Editors • Budi Purnama, Dewanta Arya Nugraha and A. Suparmi



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Preface: The 7th International Conference on Science and Applied Science (ICSAS 2022)

The 7th International Conference on Science and Applied Science (ICSAS 2022) was the sixth conference which was organized by the Physics Department, Universitas Sebelas Maret. On this occasion, the ICSAS 2022 was held virtually on 31 October, 2022, due to the COVID-19 pandemic. The ICSAS 2022 conference is aimed to bring together scholars, leading researchers, and experts from diverse backgrounds and application areas in science. Special emphasis is placed on promoting interaction between the science theoretical, experimental, and other topics related to physics.

In ICSAS 2022, there are 8 parallel sessions and four keynote speakers. The keynote speakers are Assoc. Prof. Mohsen M. Farahat from Central Metallurgical Research and Development Institute, Cairo, Cairo, Egypt; Prof. M. Aziz from Tokyo University, Japan; Prof. Madya Ts. Dr. Mohd Khairul Bin Ahmad from Universiti Tun Hussein Onn Malaysia, and Dr. Eng. Hendri Widiyandari from Universitas Sebelas Maret. While for the conference participants, there is 131 participant which was submitted abstract via the conference system. Then, the 112 full papers have been submitted from the participant, and after the reviewed process, 83 papers have been presented in the ICSAS 2022. And then for the final decision, 49 papers published in AIP Conference Proceedings.

We would like to thank all of the participants attending this conference and also to the committee for their contribution to this high-level conference and its overall success. We also would like to thank the reviewers for their positive contribution to maintain the quality of the articles presented at this conference.

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Fransiska Intan Rosari ; Sudi Mungkasi



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Co-dynamics Model of the Spread of Malaria and COVID-19 with Numerical Solutions using the Third-Order and the Fourth-Order Runge-Kutta Methods

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Abstract. We describe an existing co-dynamics model of the spread of malaria and COVID-19. Mathematical model can be used to find strategies to control the spread of malaria and COVID-19, so that their spread can be minimised in a population. The mathematical model we use is of the type of the SEI (Susceptible-Exposed-Infected) model for malaria and COVID-19 co-dynamics. For this model, we have three compartments of the human population and three compartments of the vector population. The compartments are based on S (Susceptible subpopulation), E (Exposed subpopulation), and I (Infected subpopulation). Solutions to this mathematical model can be obtained using numerical methods. We use the third-order and the fourth-order Runge-Kutta methods to solve the co-dynamics model. From our simulation results, both methods perform quite well. The fourth-order Runge-Kutta method theoretically has a higher accuracy than the third-order one. Our simulations in this paper confirm that, when solving the SEI model of the spread of malaria and COVID-19, the fourth-order Runge-Kutta method performs better than the third-order Runge-Kutta method.

INTRODUCTION

Infectious diseases are still a big problem in public health because they are one of the main causes of death in the world including in Indonesia. There are many types of infectious diseases until now that require special treatment; this includes malaria and COVID-19. Although their modes of transmission are distinct, both diseases exhibit similar symptoms like fever, headache, and muscle aches. The impact of the disease is dangerous and will be fatal to humans when someone has been infected with COVID-19 and got malaria or vice versa. Malaria is a disease caused by the bite of a female *Anopheles* mosquito with the genus of parasites *Plasmodium*, where the parasite enters and settles in the human liver organs, then infects red blood cells [1]. Four parasitic species that cause malaria in humans are *Plasmodium falciparum*, *P. vivax*, *P. malariae*, and *P. ovale*. One of the most common causes of malaria and the leading cause of death is the species *Plasmodium falciparum*. Transmission and spread of malaria are not only through mosquito bites but through organ donors, blood transfusions, the operation of syringes, and infected fetuses from their mothers [2, 3]. The spread of malaria has three factors, namely humans as hosts, the presence of *Plasmodium* in the body of female mosquitoes as agents, and the environment as a place of longing and resting mosquitoes or vectors [4]. Malaria has an incubation period. At the time of the bite, a mosquito infected with the genus *Plasmodium falciparum* requires an incubation period of 9-14 days to cause the disease, while other types of *Plasmodium* require a different incubation period [5]. The causes of spread, the number of malaria cases, the incubation period of malaria, and symptoms due to malaria infection are also explained in several studies [6–9].

Furthermore, a brand-new respiratory illness that began in Wuhan, China, shocked the world at the end of December 2019 [10, 11]. This disease spread to most countries in the world in a short period, so the World Health Organization (WHO) considered this disease a global problem which came to be known as COVID-19 or

Coronavirus disease [12]. COVID-19 is an infectious disease caused by *Severe Acute Respiratory Syndrome Coronavirus 2* (SARS-CoV-2). Coronavirus causes respiratory tract infections ranging from cold cough, *Middle East Respiratory Syndrome* (MERS), and severe acute respiratory syndrome, called *Severe Acute Respiratory Syndrome* (SARS) [13, 14]. Since March 2, 2020, COVID-19 has spread in Indonesia [15]. Transmission of COVID-19 can occur directly or indirectly through saliva, respiratory tract, and droplets released when an individual infected coughs, sneezes, talks, or sings [16]. When a susceptible individual to the disease encounters an individual infected with COVID-19, the person is not immediately infected with the virus, but becomes exposed to COVID-19 and requires follow-up health checks such as the swab test. There are two possibilities for a person exposed to COVID-19, namely negative or positive to COVID-19. Data on cases of the spread of COVID-19, symptoms due to Coronavirus infection, causes of transmission, and incubation period are also explained in several studies [17–22].

The COVID-19 pandemic has made the government impose mobility restrictions so that it can have an impact on health service programs and malaria control [23]. An individual exposed to or infected with COVID-19 does not rule out the possibility of being exposed to or infected with other diseases such as malaria and vice versa [24]. These two diseases can coexist to affect and worsen human health, as in the case of co-infection, where a previously healthy young man was later diagnosed with COVID-19 and infected with malaria [25, 26]. This case includes the co-dynamics of the spread of the disease, which is increasingly widespread at any time with a fast transmission rate. Therefore, we need a way to control the spread of malaria and COVID-19, one of which is in the field of mathematics by compiling a mathematical model called the epidemic model. Mathematical modeling helps a person to be able to understand and identify the relationship between the spread of malaria and COVID-19 with various epidemiological parameters and predict the rate of spread of malaria and COVID-19 in the future, so that it does not become more widespread.

During the incubation period of malaria and COVID-19, the population into exposed subpopulations. Several studies have discussed and developed a mathematical model for the co-dynamics of the spread of malaria and COVID-19 or other diseases with various models [27–33]. In this study, we chose not to involve the presence of vaccination, so the appropriate mathematical model for the co-dynamics of this epidemic is the SEI model (Susceptible-Exposed-Infected). We solved the co-dynamics of the spread of malaria and COVID-19 by using numerical methods, namely, Runge-Kutta methods. Runge-Kutta methods are numerical methods used to solve initial value problems in linear or nonlinear ordinary differential equations by achieving the accuracy of the Taylor series approach without requiring higher derivative calculations [34, 35]. There are several types of Runge-Kutta methods, namely the first-, second-, third-, fourth-, fifth-, and higher-order Runge-Kutta methods [35]. Thus, we decided to use two types of Runge-Kutta method, namely the third-, and the fourth-order Runge-Kutta methods, to solve the SEI model on the co-dynamics problem of the spread of malaria and COVID-19, because these two methods are quite popular and known to be sufficiently accurate.

EXPERIMENTAL DETAILS

The mathematical model for the co-dynamics of the spread of malaria and COVID-19 is the model developed in [36]. We use two numerical methods to solve the SEI model. Both of them are the third-order and the fourth-order Runge-Kutta method. Then, we use the MATLAB software for numerical calculations.

SEI Model

The spread of malaria in the SEI model has two populations. There are human and mosquito (vector) populations. The human population in the SEI model for co-dynamics of the spread of malaria and COVID-19 is separated into nine groups, namely Susceptible individuals (S_h), Exposed to malaria only (E_m), Exposed to COVID-19 only (E_c), Exposed to malaria and COVID-19 (E_{mc}), Infected individuals with malaria only (I_m), Infected individuals with COVID-19 only (I_c), Infected with malaria and exposed to COVID-19 (I_{mE_c}), Infected with both malaria and COVID-19 (I_{mc}), as well as Infected with COVID-19 and exposed to malaria (I_{cE_m}). Meanwhile, the SEI model for vector populations consists of three groups, Susceptible mosquitoes (S_v), Exposed mosquitoes to the malaria parasite (E_v), and Infectious mosquitoes (I_v). In this case, the groups I_m , I_c , I_{mE_c} , and I_{cE_m} of those who have recovered will be Susceptible individuals again, meaning that an individual is to be exposed to or reinfected with malaria and COVID-19. Our study assumes that there is a group's self-protection against COVID-19. Unlike the human population, the SEI model for vector populations already infected

with *Plasmodium* will not recover but will be alive to be infectious mosquitoes or die naturally. Tchoumi et al. [36] have formed a system of non-linear differential equations to model the spread of malaria and COVID-19 as follows

$$\left. \begin{aligned}
 \frac{dS_h}{dt} &= \Lambda_h + \omega_m I_m + \omega_c I_c + \omega_{mc} I_{mc} - (\lambda_m + \lambda_c + \mu) S_h, \\
 \frac{dE_m}{dt} &= \lambda_m S_h - (\lambda_c + \phi_m + \mu) E_m, \\
 \frac{dE_c}{dt} &= \lambda_c S_h - (\lambda_m + \phi_c + \mu) E_c, \\
 \frac{dE_{mc}}{dt} &= \lambda_c E_m + \lambda_m E_c - (\phi_{mc} + \mu) E_{mc}, \\
 \frac{dI_m}{dt} &= \phi_m E_m - (\delta \lambda_c + \omega_m + \mu) I_m, \\
 \frac{dI_c}{dt} &= \phi_c E_c - (\epsilon \lambda_m + \omega_c + \mu) I_c, \\
 \frac{dI_{mE_c}}{dt} &= \delta \lambda_c I_m - (\sigma + \mu) I_{mE_c}, \\
 \frac{dI_{mc}}{dt} &= \sigma I_{mE_c} + \phi_{mc} E_{mc} + \gamma I_{cE_m} - (\omega_{mc} + \mu) I_{mc}, \\
 \frac{dI_{cE_m}}{dt} &= \epsilon \lambda_m I_c - (\gamma + \mu) I_{cE_m}, \\
 \frac{dS_v}{dt} &= \Lambda_v - (\lambda_v + \mu_v) S_v, \\
 \frac{dE_v}{dt} &= \lambda_v S_v - (\phi_v + \mu_v) E_v, \\
 \frac{dI_v}{dt} &= \phi_v E_v - (\mu_v) I_v,
 \end{aligned} \right\} \quad (1)$$

with initial conditions

$$\begin{aligned}
 S_h(0) \geq 0, E_m(0) \geq 0, E_c(0) \geq 0, E_{mc}(0) \geq 0, I_m(0) \geq 0, \\
 I_c(0) \geq 0, I_{mE_c}(0) \geq 0, I_{mc}(0) \geq 0, I_{cE_m}(0) \geq 0.
 \end{aligned} \quad (2)$$

Equation system (1) corresponds to the schematic diagram illustrated by Tchoumi et al. [36]

In addition, the process of transmitting malaria in the mosquito (vector) population is shown as compartment diagram for the SEI model's mosquito component figure by Tchoumi et al. [36]

The description of the variables of equation system (1) for human populations (host) and mosquito populations (vector) from the SEI model is listed in Table 1.

TABLE 1. Variables of the SEI model

Variables	Description
Human (host):	
N_h	Total human population
S_h	Susceptible human subpopulation
E_m	Exposed individuals to malaria only
E_c	Exposed individuals to COVID-19 only
E_{mc}	Exposed individuals to malaria and COVID-19
I_m	Infected individuals to malaria only
I_c	Infected individuals to COVID-19 only
I_{mE_c}	Infected individuals with malaria and exposed to COVID-19
I_{mc}	Infected individuals to malaria and COVID-19
I_{cE_m}	Infected individuals with COVID-19 and exposed to malaria
Mosquitoes (vector):	
N_v	Total vector population
S_v	Susceptible mosquito subpopulation
E_v	Exposed mosquitoes to the malaria parasite
I_v	Infected mosquitoes with malaria

TABLE 2. Parameters of the SEI model

Parameters	Description
Human:	
Λ_h	Host population recruitment rate
ϕ_m	Transfer rate from exposed to infectious malaria state
ϕ_c	Transfer rate from exposed to infectious COVID-19 state
ϕ_{mc}	Proportion of individuals transferring to the co-infection I_{mc} class
ω_m	Rate of recovery of the malaria infected individuals
ω_c	Rate of recovery of the COVID-19 infected individuals
ω_{mc}	Rate of recovery of the malaria and COVID-19 infectious individuals
μ	Host population death rate
δ	Enhancement factor of acquiring COVID-19 following malaria infection
ϵ	Enhancement factor of acquiring malaria following COVID-19 infection
σ	Infection rate of COVID-19 of individuals already infected with malaria
γ	Malaria infection rate of individuals already infected with COVID-19
κ	Proportion of individuals employing personal protection
ζ	Efficacy of personal protection
b	Number of female mosquito bites per day
β_m	Malaria transmission probability per mosquito bite
β_c	Transmission probability of COVID-19 per contact
λ_m	Forces of infection from malaria
λ_c	Forces of infection from COVID-19
Mosquitoes:	
Λ_v	Vector recruitment rate
ϕ_m	Transfer rate from exposed to infectious class
μ_v	Vector natural death rate
β_v	Transmission probability in vector from infected humans
λ_v	Forces of infection from a bite of the malaria parasite

Furthermore, the description of the parameters of equation (1) for human populations and mosquito populations (vector) from the SEI model is listed in Table 2.

In solving the SEI model of equation system (1) and both compartment diagrams for the co-dynamics of the spread of malaria and COVID-19, we choose two methods to obtain solutions, namely the third-, and the fourth-order Runge-Kutta methods.

Third-Order Runge-Kutta Method

Consider the initial value problem $dy/dx = f(x, y)$, $y(x_0) = x_0$. Triatmodjo [37] wrote the scheme for the third-order Runge-Kutta method as follows

$$y_{i+1} = y_i + \frac{1}{6}(k_1 + 4k_2 + k_3)h, \quad (3)$$

where h is the step size of the free variable and

$$\begin{aligned} k_1 &= f(x_i, y_i), \\ k_2 &= f\left(x_i + \frac{1}{2}h, y_i + \frac{1}{2}k_1h\right), \\ k_3 &= f(x_i + h, y_i - k_1h + 2k_2h). \end{aligned} \quad (4)$$

We take into account the initial value of each subpopulation $S_{h(1)}, E_{m(1)}, E_{c(1)}, E_{mc(1)}, I_{m(1)}, I_{c(1)}, I_{mE_c(1)}, I_{mc(1)}, I_{cEm(1)}, S_{v(1)}, E_{v(1)}, I_{v(1)}$, and $N_{h(1)}$ from $S_h(0), E_m(0), E_c(0), E_{mc}(0), I_m(0), I_c(0), I_{mE_c}(0), I_{mc}(0), I_{cEm}(0), S_v(0), E_v(0), I_v(0)$, and $N_h(0)$, respectively, the third-order Runge-Kutta method to solve the SEI model (1) is

$$\begin{aligned} S_{h(i+1)} &= S_{h(i)} + \frac{1}{6}(k_{1S_h} + 4k_{2S_h} + k_{3S_h})h, \\ E_{m(i+1)} &= E_{m(i)} + \frac{1}{6}(k_{1E_m} + 4k_{2E_m} + k_{3E_m})h, \\ E_{c(i+1)} &= E_{c(i)} + \frac{1}{6}(k_{1E_c} + 4k_{2E_c} + k_{3E_c})h, \\ E_{mc(i+1)} &= E_{mc(i)} + \frac{1}{6}(k_{1E_{mc}} + 4k_{2E_{mc}} + k_{3E_{mc}})h, \\ I_{m(i+1)} &= I_{m(i)} + \frac{1}{6}(k_{1I_m} + 4k_{2I_m} + k_{3I_m})h, \\ I_{c(i+1)} &= I_{c(i)} + \frac{1}{6}(k_{1I_c} + 4k_{2I_c} + k_{3I_c})h, \\ I_{mE_c(i+1)} &= I_{mE_c(i)} + \frac{1}{6}(k_{1I_{mE_c}} + 4k_{2I_{mE_c}} + k_{3I_{mE_c}})h, \\ I_{mc(i+1)} &= I_{mc(i)} + \frac{1}{6}(k_{1I_{mc}} + 4k_{2I_{mc}} + k_{3I_{mc}})h, \\ I_{cEm(i+1)} &= I_{cEm(i)} + \frac{1}{6}(k_{1I_{cEm}} + 4k_{2I_{cEm}} + k_{3I_{cEm}})h, \\ S_{v(i+1)} &= S_{v(i)} + \frac{1}{6}(k_{1S_v} + 4k_{2S_v} + k_{3S_v})h, \\ E_{v(i+1)} &= E_{v(i)} + \frac{1}{6}(k_{1E_v} + 4k_{2E_v} + k_{3E_v})h, \\ I_{v(i+1)} &= I_{v(i)} + \frac{1}{6}(k_{1I_v} + 4k_{2I_v} + k_{3I_v})h, \end{aligned} \quad (5)$$

$$N_{h(i+1)} = S_{h(i+1)} + E_{m(i+1)} + E_{c(i+1)} + E_{mc(i+1)} + I_{m(i+1)} + I_{c(i+1)} + I_{mE_c(i+1)} + I_{mc(i+1)} + I_{cEm(i+1)},$$

$$\lambda_c = \beta_c(1 - \kappa\zeta) \frac{(I_{c(i+1)} + I_{mc(i+1)} + I_{cEm(i+1)})}{N_{h(i+1)}},$$

$$\lambda_m = \beta_m b \frac{I_{v(i+1)}}{N_{h(i+1)}},$$

$$\lambda_v = \beta_v b \frac{(I_{m(i+1)} + I_{mc(i+1)} + I_{mE_c(i+1)})}{N_{h(i+1)}}.$$

Here, h in the time step, i is the iteration index and:

$$\begin{aligned}
k_{1S_h} &= \Lambda_h + \omega_m I_{m(i)} + \omega_c I_{c(i)} + \omega_{mc} I_{mc(i)} - (\lambda_m + \lambda_c + \mu) S_{h(i)}, \\
k_{1E_m} &= \lambda_m S_{h(i)} - (\lambda_c + \phi_m + \mu) E_{m(i)}, \\
k_{1E_c} &= \lambda_c S_{h(i)} - (\lambda_m + \phi_c + \mu) E_{c(i)}, \\
k_{1E_{mc}} &= \lambda_c E_{m(i)} + \lambda_m E_{c(i)} - (\phi_{mc} + \mu) E_{mc(i)}, \\
k_{1I_m} &= \phi_m E_{m(i)} - (\delta\lambda_c + \omega_m + \mu) I_{m(i)}, \\
k_{1I_c} &= \phi_c E_{c(i)} - (\epsilon\lambda_m + \omega_c + \mu) I_{c(i)}, \\
k_{1I_{mE_c}} &= \delta\lambda_c I_{m(i)} - (\sigma + \mu) I_{mE_c(i)}, \\
k_{1I_{mc}} &= \sigma I_{mE_c(i)} + \phi_{mc} E_{mc(i)} + \gamma I_{cE_m(i)} - (\omega_{mc} + \mu) I_{mc(i)}, \\
k_{1I_{cE_m}} &= \epsilon\lambda_m I_{c(i)} - (\gamma + \mu) I_{cE_m(i)}, \\
k_{1S_v} &= \Lambda_v - (\lambda_v + \mu_v) S_{v(i)}, \\
k_{1E_v} &= \lambda_v S_{v(i)} - (\phi_v + \mu_v) E_{v(i)}, \\
k_{1I_v} &= \phi_v E_{v(i)} - (\mu_v) I_{v(i)},
\end{aligned}$$

$$\begin{aligned}
k_{2N_h} &= S_{h(i)} + \frac{1}{2}k_{1S_h}h + E_{m(i)} + \frac{1}{2}k_{1E_m}h + E_{c(i)} + \frac{1}{2}k_{1E_c}h + E_{mc(i)} + \frac{1}{2}k_{1E_{mc}}h + I_{m(i)} + \frac{1}{2}k_{1I_m}h \\
&\quad + I_{c(i)} + \frac{1}{2}k_{1I_c}h + I_{mE_c(i)} + \frac{1}{2}k_{1I_{mE_c}}h + I_{mc(i)} + \frac{1}{2}k_{1I_{mc}}h + I_{cE_m(i)} + \frac{1}{2}k_{1I_{cE_m}}h, \\
\lambda_{c2} &= \beta_c(1 - \kappa\zeta) \frac{\left(I_{c(i)} + \frac{1}{2}k_{1I_c}h + I_{mc(i)} + \frac{1}{2}k_{1I_{mc}}h + I_{cE_m(i)} + \frac{1}{2}k_{1I_{cE_m}}h \right)}{k_{2N_h}},
\end{aligned}$$

$$\lambda_{m2} = \beta_m b \frac{\left(I_{v(i)} + \frac{1}{2}k_{1I_v}h \right)}{k_{2N_h}},$$

$$\lambda_{v2} = \beta_v b \frac{\left(I_{m(i)} + \frac{1}{2}k_{1I_m}h + I_{mc(i)} + \frac{1}{2}k_{1I_{mc}}h + I_{mE_c(i)} + \frac{1}{2}k_{1I_{mE_c}}h \right)}{k_{2N_h}},$$

(6)

$$\begin{aligned}
k_{2S_h} &= \Lambda_h + \omega_m \left(I_{m(i)} + \frac{1}{2}k_{1I_m}h \right) + \omega_c \left(I_{c(i)} + \frac{1}{2}k_{1I_c}h \right) + \omega_{mc} \left(I_{mc(i)} + \frac{1}{2}k_{1I_{mc}}h \right) - (\lambda_{m2} + \lambda_{c2} + \mu) \\
&\quad \left(S_{h(i)} + \frac{1}{2}k_{1S_h}h \right),
\end{aligned}$$

$$k_{2E_m} = \lambda_{m2} \left(S_{h(i)} + \frac{1}{2}k_{1S_h}h \right) - (\lambda_{c2} + \phi_m + \mu) \left(E_{m(i)} + \frac{1}{2}k_{1E_m}h \right),$$

$$k_{2E_c} = \lambda_{c2} \left(S_{h(i)} + \frac{1}{2}k_{1S_h}h \right) - (\lambda_{m2} + \phi_c + \mu) \left(E_{c(i)} + \frac{1}{2}k_{1E_c}h \right),$$

$$k_{2E_{mc}} = \lambda_{c2} \left(E_{m(i)} + \frac{1}{2}k_{1E_m}h \right) + \lambda_{m2} \left(E_{c(i)} + \frac{1}{2}k_{1E_c}h \right) - (\phi_{mc} + \mu) \left(E_{mc(i)} + \frac{1}{2}k_{1E_{mc}}h \right),$$

$$k_{2I_m} = \phi_m \left(E_{m(i)} + \frac{1}{2}k_{1E_m}h \right) - (\delta\lambda_{c2} + \omega_m + \mu) \left(I_{m(i)} + \frac{1}{2}k_{1I_m}h \right),$$

$$k_{2I_c} = \phi_c \left(E_{c(i)} + \frac{1}{2}k_{1E_c}h \right) - (\epsilon\lambda_{m2} + \omega_c + \mu) \left(I_{c(i)} + \frac{1}{2}k_{1I_c}h \right),$$

$$k_{2I_{mE_c}} = \delta\lambda_{c2} \left(I_{m(i)} + \frac{1}{2}k_{1I_m}h \right) - (\sigma + \mu) \left(I_{mE_c(i)} + \frac{1}{2}k_{1I_{mE_c}}h \right),$$

$$\begin{aligned}
k_{2I_{mc}} &= \sigma \left(I_{mE_c(i)} + \frac{1}{2}k_{1I_{mE_c}}h \right) + \phi_{mc} \left(E_{mc(i)} + \frac{1}{2}k_{1E_{mc}}h \right) + \gamma \left(I_{cE_m(i)} + \frac{1}{2}k_{1I_{cE_m}}h \right) - (\omega_{mc} + \mu) \\
&\quad \left(I_{mc(i)} + \frac{1}{2}k_{1I_{mc}}h \right),
\end{aligned}$$

$$k_{2I_{cE_m}} = \epsilon\lambda_{m2} \left(I_{c(i)} + \frac{1}{2}k_{1I_c}h \right) - (\gamma + \mu) \left(I_{cE_m(i)} + \frac{1}{2}k_{1I_{cE_m}}h \right),$$

$$k_{2S_v} = \Lambda_v - (\lambda_{v2} + \mu_v) \left(S_{v(i)} + \frac{1}{2}k_{1S_v}h \right),$$

$$k_{2E_v} = \lambda_{v2} \left(S_{v(i)} + \frac{1}{2}k_{1S_v}h \right) - (\phi_v + \mu_v) \left(E_{v(i)} + \frac{1}{2}k_{1E_v}h \right),$$

$$k_{2I_v} = \phi_v \left(E_{v(i)} + \frac{1}{2} k_{1E_v} h \right) - (\mu_v) \left(I_{v(i)} + \frac{1}{2} k_{1I_v} h \right),$$

$$k_{3N_h} = (S_{h(i)} - k_{1S_h} h + 2k_{2S_h} h) + (E_{m(i)} - k_{1E_m} h + 2k_{2E_m} h) + (E_{c(i)} - k_{1E_c} h + 2k_{2E_c} h) \\ + (E_{mc(i)} - k_{1E_{mc}} h + 2k_{2E_{mc}} h) + (I_{m(i)} - k_{1I_m} h + 2k_{2I_m} h) + (I_{c(i)} - k_{1I_c} h + 2k_{2I_c} h) \\ + (I_{mE_c(i)} - k_{1I_{mE_c}} h + 2k_{2I_{mE_c}} h) + (I_{mc(i)} - k_{1I_{mc}} h + 2k_{2I_{mc}} h) \\ + (I_{cE_m(i)} - k_{1I_{cE_m}} h + 2k_{2I_{cE_m}} h),$$

$$\lambda_{c3} = \beta_c (1 - \kappa \zeta) \frac{\left((I_{c(i)} - k_{1I_c} h + 2k_{2I_c} h) + (I_{mc(i)} - k_{1I_{mc}} h + 2k_{2I_{mc}} h) \right) \\ + (I_{cE_m(i)} - k_{1I_{cE_m}} h + 2k_{2I_{cE_m}} h)}{k_{3N_h}},$$

$$\lambda_{m3} = \beta_m b \frac{(I_{v(i)} - k_{1I_v} h + 2k_{2I_v} h)}{k_{3N_h}},$$

$$\lambda_{v3} = \beta_v b \frac{\left((I_{m(i)} - k_{1I_m} h + 2k_{2I_m} h) + (I_{mc(i)} - k_{1I_{mc}} h + 2k_{2I_{mc}} h) \right) \\ + (I_{mE_c(i)} - k_{1I_{mE_c}} h + 2k_{2I_{mE_c}} h)}{k_{3N_h}},$$

$$k_{3S_h} = \Lambda_h + \omega_m (I_{m(i)} - k_{1I_m} h + 2k_{2I_m} h) + \omega_c (I_{c(i)} - k_{1I_c} h + 2k_{2I_c} h) \\ + \omega_{mc} (I_{mc(i)} - k_{1I_{mc}} h + 2k_{2I_{mc}} h) - (\lambda_{m3} + \lambda_{c3} + \mu) (S_{h(i)} - k_{1S_h} h + 2k_{2S_h} h),$$

$$k_{3E_m} = \lambda_{m3} (S_{h(i)} - k_{1S_h} h + 2k_{2S_h} h) - (\lambda_{c3} + \phi_m + \mu) (E_{m(i)} - k_{1E_m} h + 2k_{2E_m} h),$$

$$k_{3E_c} = \lambda_{c3} (S_{h(i)} - k_{1S_h} h + 2k_{2S_h} h) - (\lambda_{m3} + \phi_c + \mu) (E_{c(i)} - k_{1E_c} h + 2k_{2E_c} h),$$

$$k_{3E_{mc}} = \lambda_{c3} (E_{m(i)} - k_{1E_m} h + 2k_{2E_m} h) + \lambda_{m3} (E_{c(i)} - k_{1E_c} h + 2k_{2E_c} h) - (\phi_{mc} + \mu) \\ (E_{mc(i)} - k_{1E_{mc}} h + 2k_{2E_{mc}} h),$$

$$k_{3I_m} = \phi_m (E_{m(i)} - k_{1E_m} h + 2k_{2E_m} h) - (\delta \lambda_{c3} + \omega_m + \mu) (I_{m(i)} - k_{1I_m} h + 2k_{2I_m} h),$$

$$k_{3I_c} = \phi_c (E_{c(i)} - k_{1E_c} h + 2k_{2E_c} h) - (\epsilon \lambda_{m3} + \omega_c + \mu) (I_{c(i)} - k_{1I_c} h + 2k_{2I_c} h),$$

$$k_{3I_{mE_c}} = \delta \lambda_{c3} (I_{m(i)} - k_{1I_m} h + 2k_{2I_m} h) - (\sigma + \mu) (I_{mE_c(i)} - k_{1I_{mE_c}} h + 2k_{2I_{mE_c}} h),$$

$$k_{3I_{mc}} = \sigma (I_{mE_c(i)} - k_{1I_{mE_c}} h + 2k_{2I_{mE_c}} h) + \phi_{mc} (E_{mc(i)} - k_{1E_{mc}} h + 2k_{2E_{mc}} h) \\ + \gamma (I_{cE_m(i)} - k_{1I_{cE_m}} h + 2k_{2I_{cE_m}} h) - (\omega_{mc} + \mu) (I_{mc(i)} - k_{1I_{mc}} h + 2k_{2I_{mc}} h),$$

$$k_{3I_{cE_m}} = \epsilon \lambda_{m3} (I_{c(i)} - k_{1I_c} h + 2k_{2I_c} h) - (\gamma + \mu) (I_{cE_m(i)} - k_{1I_{cE_m}} h + 2k_{2I_{cE_m}} h),$$

$$k_{3S_v} = \Lambda_v - (\lambda_{v3} + \mu_v) (S_{v(i)} - k_{1S_v} h + 2k_{2S_v} h),$$

$$k_{3E_v} = \lambda_{v3} (S_{v(i)} - k_{1S_v} h + 2k_{2S_v} h) - (\phi_v + \mu_v) (E_{v(i)} - k_{1E_v} h + 2k_{2E_v} h),$$

$$k_{3I_v} = \phi_v (E_{v(i)} - k_{1E_v} h + 2k_{2E_v} h) - (\mu_v) (I_{v(i)} - k_{1I_v} h + 2k_{2I_v} h).$$

Fourth-Order Runge-Kutta Method

The fourth-order Runge-Kutta method is one of the numerical methods for solving differential equations that can be derived from Taylor expansions [38, 39]. This method has a higher order of solution accuracy than the first-, the second-, and the third-order methods [40]. Reconsider the initial value problem $dy/dx = f(x, y)$, $y(x_0) = x_0$, Chapra and Canale [41] wrote down the most commonly used form of the classical fourth-order Runge-Kutta method as follows

$$y_{i+1} = y_i + \frac{1}{6} (k_1 + 2k_2 + 2k_3 + k_4)h, \quad (7)$$

where

$$\begin{aligned}
k_1 &= f(x_i, y_i), \\
k_2 &= f\left(x_i + \frac{1}{2}h, y_i + \frac{1}{2}k_1h\right), \\
k_3 &= f\left(x_i + \frac{1}{2}h, y_i + \frac{1}{2}k_2h\right), \\
k_4 &= f(x_i + h, y_i + k_3h).
\end{aligned} \tag{8}$$

Now, we take into account the initial value of each subpopulation $S_{h(1)}, E_{m(1)}, E_{c(1)}, E_{mc(1)}, I_{m(1)}, I_{c(1)}, I_{mE_c(1)}, I_{mc(1)}, I_{cEm(1)}, S_{v(1)}, E_{v(1)}, I_{v(1)}$, and $N_{h(1)}$ from $S_h(0), E_m(0), E_c(0), E_{mc}(0), I_m(0), I_c(0), I_{mE_c}(0), I_{mc}(0), I_{cEm}(0), S_v(0), E_v(0), I_v(0)$, and $N_h(0)$, respectively, the fourth-order Runge-Kutta method to solve the SEI model (1) is

$$\begin{aligned}
S_{h(i+1)} &= S_{h(i)} + \frac{1}{6}(k_{1S_h} + 2k_{2S_h} + 2k_{3S_h} + k_{4S_h})h, \\
E_{m(i+1)} &= E_{m(i)} + \frac{1}{6}(k_{1E_m} + 2k_{2E_m} + 2k_{3E_m} + k_{4E_m})h, \\
E_{c(i+1)} &= E_{c(i)} + \frac{1}{6}(k_{1E_c} + 2k_{2E_c} + 2k_{3E_c} + k_{4E_c})h, \\
E_{mc(i+1)} &= E_{mc(i)} + \frac{1}{6}(k_{1E_{mc}} + 2k_{2E_{mc}} + 2k_{3E_{mc}} + k_{4E_{mc}})h, \\
I_{m(i+1)} &= I_{m(i)} + \frac{1}{6}(k_{1I_m} + 2k_{2I_m} + 2k_{3I_m} + k_{4I_m})h, \\
I_{c(i+1)} &= I_{c(i)} + \frac{1}{6}(k_{1I_c} + 2k_{2I_c} + 2k_{3I_c} + k_{4I_c})h, \\
I_{mE_c(i+1)} &= I_{mE_c(i)} + \frac{1}{6}(k_{1I_{mE_c}} + 2k_{2I_{mE_c}} + 2k_{3I_{mE_c}} + k_{4I_{mE_c}})h, \\
I_{mc(i+1)} &= I_{mc(i)} + \frac{1}{6}(k_{1I_{mc}} + 2k_{2I_{mc}} + 2k_{3I_{mc}} + k_{4I_{mc}})h, \\
I_{cEm(i+1)} &= I_{cEm(i)} + \frac{1}{6}(k_{1I_{cEm}} + 2k_{2I_{cEm}} + 2k_{3I_{cEm}} + k_{4I_{cEm}})h, \\
S_{v(i+1)} &= S_{v(i)} + \frac{1}{6}(k_{1S_v} + 2k_{2S_v} + 2k_{3S_v} + k_{4S_v})h, \\
E_{v(i+1)} &= E_{v(i)} + \frac{1}{6}(k_{1E_v} + 2k_{2E_v} + 2k_{3E_v} + k_{4E_v})h, \\
I_{v(i+1)} &= I_{v(i)} + \frac{1}{6}(k_{1I_v} + 2k_{2I_v} + 2k_{3I_v} + k_{4I_v})h, \\
N_{h(i+1)} &= S_{h(i+1)} + E_{m(i+1)} + E_{c(i+1)} + E_{mc(i+1)} + I_{m(i+1)} + I_{c(i+1)} + I_{mE_c(i+1)} + I_{mc(i+1)} + I_{cEm(i+1)} \\
\lambda_c &= \beta_c(1 - \kappa\zeta) \frac{(I_{c(i+1)} + I_{mc(i+1)} + I_{cEm(i+1)})}{N_{h(i+1)}}, \\
\lambda_m &= \beta_m b \frac{I_{v(i+1)}}{N_{h(i+1)}}, \\
\lambda_v &= \beta_v b \frac{(I_{m(i+1)} + I_{mc(i+1)} + I_{mE_c(i+1)})}{N_{h(i+1)}}.
\end{aligned} \tag{9}$$

Here, h is the time step, i is the iteration index and:

$$\begin{aligned}
k_{1S_h} &= \Lambda_h + \omega_m I_{m(i)} + \omega_c I_{c(i)} + \omega_{mc} I_{mc(i)} - (\lambda_m + \lambda_c + \mu) S_{h(i)}, \\
k_{1E_m} &= \lambda_m S_{h(i)} - (\lambda_c + \phi_m + \mu) E_{m(i)}, \\
k_{1E_c} &= \lambda_c S_{h(i)} - (\lambda_m + \phi_c + \mu) E_{c(i)}, \\
k_{1E_{mc}} &= \lambda_c E_{m(i)} + \lambda_m E_{c(i)} - (\phi_{mc} + \mu) E_{mc(i)}, \\
k_{1I_m} &= \phi_m E_{m(i)} - (\delta\lambda_c + \omega_m + \mu) I_{m(i)}, \\
k_{1I_c} &= \phi_c E_{c(i)} - (\epsilon\lambda_m + \omega_c + \mu) I_{c(i)}, \\
k_{1I_{mE_c}} &= \delta\lambda_c I_{m(i)} - (\sigma + \mu) I_{mE_c(i)}, \\
k_{1I_{mc}} &= \sigma I_{mE_c(i)} + \phi_{mc} E_{mc(i)} + \gamma I_{cEm(i)} - (\omega_{mc} + \mu) I_{mc(i)},
\end{aligned} \tag{10}$$

$$\begin{aligned}
k_{1cEm} &= \epsilon \lambda_m I_{c(i)} - (\gamma + \mu) I_{cEm(i)}, \\
k_{1Sv} &= \Lambda_v - (\lambda_v + \mu_v) S_{v(i)}, \\
k_{1Ev} &= \lambda_v S_{v(i)} - (\phi_v + \mu_v) E_{v(i)}, \\
k_{1Iv} &= \phi_v E_{v(i)} - (\mu_v) I_{v(i)},
\end{aligned}$$

$$\begin{aligned}
k_{2N_h} &= S_{h(i)} + \frac{1}{2} k_{1S_h} h + E_{m(i)} + \frac{1}{2} k_{1E_m} h + E_{c(i)} + \frac{1}{2} k_{1E_c} h + E_{mc(i)} + \frac{1}{2} k_{1Emc} h + I_{m(i)} + \frac{1}{2} k_{1I_m} h \\
&\quad + I_{c(i)} + \frac{1}{2} k_{1I_c} h + I_{mE_c(i)} + \frac{1}{2} k_{1ImE_c} h + I_{mc(i)} + \frac{1}{2} k_{1Imc} h + I_{cEm(i)} + \frac{1}{2} k_{1IcEm} h,
\end{aligned}$$

$$\lambda_{c2} = \beta_c (1 - \kappa \zeta) \frac{\left(I_{c(i)} + \frac{1}{2} k_{1I_c} h + I_{mc(i)} + \frac{1}{2} k_{1Imc} h + I_{cEm(i)} + \frac{1}{2} k_{1IcEm} h \right)}{k_{2N_h}},$$

$$\lambda_{m2} = \beta_m b \frac{\left(I_{v(i)} + \frac{1}{2} k_{1I_v} h \right)}{k_{2N_h}},$$

$$\lambda_{v2} = \beta_v b \frac{\left(I_{m(i)} + \frac{1}{2} k_{1I_m} h + I_{mc(i)} + \frac{1}{2} k_{1Imc} h + I_{mE_c(i)} + \frac{1}{2} k_{1ImE_c} h \right)}{k_{2N_h}},$$

$$\begin{aligned}
k_{2S_h} &= \Lambda_h + \omega_m \left(I_{m(i)} + \frac{1}{2} k_{1I_m} h \right) + \omega_c \left(I_{c(i)} + \frac{1}{2} k_{1I_c} h \right) + \omega_{mc} \left(I_{mc(i)} + \frac{1}{2} k_{1Imc} h \right) - (\lambda_{m2} + \lambda_{c2} + \mu) \\
&\quad \left(S_{h(i)} + \frac{1}{2} k_{1S_h} h \right),
\end{aligned}$$

$$k_{2E_m} = \lambda_{m2} \left(S_{h(i)} + \frac{1}{2} k_{1S_h} h \right) - (\lambda_{c2} + \phi_m + \mu) \left(E_{m(i)} + \frac{1}{2} k_{1E_m} h \right),$$

$$k_{2E_c} = \lambda_{c2} \left(S_{h(i)} + \frac{1}{2} k_{1S_h} h \right) - (\lambda_{m2} + \phi_c + \mu) \left(E_{c(i)} + \frac{1}{2} k_{1E_c} h \right),$$

$$k_{2Emc} = \lambda_{c2} \left(E_{m(i)} + \frac{1}{2} k_{1E_m} h \right) + \lambda_{m2} \left(E_{c(i)} + \frac{1}{2} k_{1E_c} h \right) - (\phi_{mc} + \mu) \left(E_{mc(i)} + \frac{1}{2} k_{1Emc} h \right),$$

$$k_{2I_m} = \phi_m \left(E_{m(i)} + \frac{1}{2} k_{1E_m} h \right) - (\delta \lambda_{c2} + \omega_m + \mu) \left(I_{m(i)} + \frac{1}{2} k_{1I_m} h \right),$$

$$k_{2I_c} = \phi_c \left(E_{c(i)} + \frac{1}{2} k_{1E_c} h \right) - (\epsilon \lambda_{m2} + \omega_c + \mu) \left(I_{c(i)} + \frac{1}{2} k_{1I_c} h \right),$$

$$k_{2ImE_c} = \delta \lambda_{c2} \left(I_{m(i)} + \frac{1}{2} k_{1I_m} h \right) - (\sigma + \mu) \left(I_{mE_c(i)} + \frac{1}{2} k_{1ImE_c} h \right),$$

$$\begin{aligned}
k_{2Imc} &= \sigma \left(I_{mE_c(i)} + \frac{1}{2} k_{1ImE_c} h \right) + \phi_{mc} \left(E_{mc(i)} + \frac{1}{2} k_{1Emc} h \right) + \gamma \left(I_{cEm(i)} + \frac{1}{2} k_{1IcEm} h \right) - (\omega_{mc} + \mu) \\
&\quad \left(I_{mc(i)} + \frac{1}{2} k_{1Imc} h \right),
\end{aligned}$$

$$k_{2IcEm} = \epsilon \lambda_{m2} \left(I_{c(i)} + \frac{1}{2} k_{1I_c} h \right) - (\gamma + \mu) \left(I_{cEm(i)} + \frac{1}{2} k_{1IcEm} h \right),$$

$$k_{2S_v} = \Lambda_v - (\lambda_{v2} + \mu_v) \left(S_{v(i)} + \frac{1}{2} k_{1S_v} h \right),$$

$$k_{2E_v} = \lambda_{v2} \left(S_{v(i)} + \frac{1}{2} k_{1S_v} h \right) - (\phi_v + \mu_v) \left(E_{v(i)} + \frac{1}{2} k_{1E_v} h \right),$$

$$k_{2I_v} = \phi_v \left(E_{v(i)} + \frac{1}{2} k_{1E_v} h \right) - (\mu_v) \left(I_{v(i)} + \frac{1}{2} k_{1I_v} h \right),$$

$$\begin{aligned}
k_{3N_h} &= S_{h(i)} + \frac{1}{2} k_{2S_h} h + E_{m(i)} + \frac{1}{2} k_{2E_m} h + E_{c(i)} + \frac{1}{2} k_{2E_c} h + E_{mc(i)} + \frac{1}{2} k_{2Emc} h + I_{m(i)} + \frac{1}{2} k_{2I_m} h \\
&\quad + I_{c(i)} + \frac{1}{2} k_{2I_c} h + I_{mE_c(i)} + \frac{1}{2} k_{2ImE_c} h + I_{mc(i)} + \frac{1}{2} k_{2Imc} h + I_{cEm(i)} + \frac{1}{2} k_{2IcEm} h,
\end{aligned}$$

$$\lambda_{c3} = \beta_c (1 - \kappa \zeta) \frac{\left(I_{c(i)} + \frac{1}{2} k_{2I_c} h + I_{mc(i)} + \frac{1}{2} k_{2Imc} h + I_{cEm(i)} + \frac{1}{2} k_{2IcEm} h \right)}{k_{3N_h}},$$

$$\begin{aligned}
\lambda_{m3} &= \beta_m b \frac{\left(I_{v(i)} + \frac{1}{2}k_{2I_v}h\right)}{k_{3N_h}}, \\
\lambda_{v3} &= \beta_v b \frac{\left(I_{m(i)} + \frac{1}{2}k_{2I_m}h + I_{mc(i)} + \frac{1}{2}k_{2I_{mc}}h + I_{mE_c(i)} + \frac{1}{2}k_{2I_{mE_c}}h\right)}{k_{3N_h}}, \\
k_{3S_h} &= \Lambda_h + \omega_m \left(I_{m(i)} + \frac{1}{2}k_{2I_m}h\right) + \omega_c \left(I_{c(i)} + \frac{1}{2}k_{2I_c}h\right) + \omega_{mc} \left(I_{mc(i)} + \frac{1}{2}k_{2I_{mc}}h\right) - (\lambda_{m3} + \lambda_{c3} + \mu) \\
&\quad \left(S_{h(i)} + \frac{1}{2}k_{2S_h}h\right), \\
k_{3E_m} &= \lambda_{m3} \left(S_{h(i)} + \frac{1}{2}k_{2S_h}h\right) - (\lambda_{c3} + \phi_m + \mu) \left(E_{m(i)} + \frac{1}{2}k_{2E_m}h\right), \\
k_{3E_c} &= \lambda_{c3} \left(S_{h(i)} + \frac{1}{2}k_{2S_h}h\right) - (\lambda_{m3} + \phi_c + \mu) \left(E_{c(i)} + \frac{1}{2}k_{2E_c}h\right), \\
k_{3E_{mc}} &= \lambda_{c3} \left(E_{m(i)} + \frac{1}{2}k_{2E_m}h\right) + \lambda_{m3} \left(E_{c(i)} + \frac{1}{2}k_{2E_c}h\right) - (\phi_{mc} + \mu) \left(E_{mc(i)} + \frac{1}{2}k_{2E_{mc}}h\right), \\
k_{3I_m} &= \phi_m \left(E_{m(i)} + \frac{1}{2}k_{2E_m}h\right) - (\delta\lambda_{c3} + \omega_m + \mu) \left(I_{m(i)} + \frac{1}{2}k_{2I_m}h\right), \\
k_{3I_c} &= \phi_c \left(E_{c(i)} + \frac{1}{2}k_{2E_c}h\right) - (\epsilon\lambda_{m3} + \omega_c + \mu) \left(I_{c(i)} + \frac{1}{2}k_{2I_c}h\right), \\
k_{3I_{mE_c}} &= \delta\lambda_{c3} \left(I_{m(i)} + \frac{1}{2}k_{2I_m}h\right) - (\sigma + \mu) \left(I_{mE_c(i)} + \frac{1}{2}k_{2I_{mE_c}}h\right), \\
k_{3I_{mc}} &= \sigma \left(I_{mE_c(i)} + \frac{1}{2}k_{2I_{mE_c}}h\right) + \phi_{mc} \left(E_{mc(i)} + \frac{1}{2}k_{2E_{mc}}h\right) + \gamma \left(I_{cE_m(i)} + \frac{1}{2}k_{2I_{cE_m}}h\right) - (\omega_{mc} + \mu) \\
&\quad \left(I_{mc(i)} + \frac{1}{2}k_{2I_{mc}}h\right), \\
k_{3I_{cE_m}} &= \epsilon\lambda_{m3} \left(I_{c(i)} + \frac{1}{2}k_{2I_c}h\right) - (\gamma + \mu) \left(I_{cE_m(i)} + \frac{1}{2}k_{2I_{cE_m}}h\right), \\
k_{3S_v} &= \Lambda_v - (\lambda_{v3} + \mu_v) \left(S_{v(i)} + \frac{1}{2}k_{2S_v}h\right), \\
k_{3E_v} &= \lambda_{v3} \left(S_{v(i)} + \frac{1}{2}k_{2S_v}h\right) - (\phi_v + \mu_v) \left(E_{v(i)} + \frac{1}{2}k_{2E_v}h\right), \\
k_{3I_v} &= \phi_v \left(E_{v(i)} + \frac{1}{2}k_{2E_v}h\right) - (\mu_v) \left(I_{v(i)} + \frac{1}{2}k_{2I_v}h\right), \\
k_{4N_h} &= S_{h(i)} + k_{3S_h}h + E_{m(i)} + k_{3E_m}h + E_{c(i)} + k_{3E_c}h + E_{mc(i)} + k_{3E_{mc}}h + I_{m(i)} + k_{3I_m}h \\
&\quad + I_{c(i)} + k_{3I_c}h + I_{mE_c(i)} + k_{3I_{mE_c}}h + I_{mc(i)} + k_{3I_{mc}}h + I_{cE_m(i)} + k_{3I_{cE_m}}h, \\
\lambda_{c4} &= \beta_c (1 - \kappa\zeta) \frac{\left(I_{c(i)} + k_{3I_c}h + I_{mc(i)} + k_{3I_{mc}}h + I_{cE_m(i)} + k_{3I_{cE_m}}h\right)}{k_{4N_h}}, \\
\lambda_{m4} &= \beta_m b \frac{\left(I_{v(i)} + k_{3I_v}h\right)}{k_{4N_h}}, \\
\lambda_{v4} &= \beta_v b \frac{\left(I_{m(i)} + k_{3I_m}h + I_{mc(i)} + k_{3I_{mc}}h + I_{mE_c(i)} + k_{3I_{mE_c}}h\right)}{k_{4N_h}}, \\
k_{4S_h} &= \Lambda_h + \omega_m \left(I_{m(i)} + k_{3I_m}h\right) + \omega_c \left(I_{c(i)} + k_{3I_c}h\right) + \omega_{mc} \left(I_{mc(i)} + k_{3I_{mc}}h\right) - (\lambda_{m4} + \lambda_{c4} + \mu) \\
&\quad \left(S_{h(i)} + k_{3S_h}h\right), \\
k_{4E_m} &= \lambda_{m4} \left(S_{h(i)} + k_{3S_h}h\right) - (\lambda_{c4} + \phi_m + \mu) \left(E_{m(i)} + k_{3E_m}h\right), \\
k_{4E_c} &= \lambda_{c4} \left(S_{h(i)} + k_{3S_h}h\right) - (\lambda_{m4} + \phi_c + \mu) \left(E_{c(i)} + k_{3E_c}h\right), \\
k_{4E_{mc}} &= \lambda_{c4} \left(E_{m(i)} + k_{3E_m}h\right) + \lambda_{m4} \left(E_{c(i)} + k_{3E_c}h\right) - (\phi_{mc} + \mu) \left(E_{mc(i)} + k_{3E_{mc}}h\right), \\
k_{4I_m} &= \phi_m \left(E_{m(i)} + k_{3E_m}h\right) - (\delta\lambda_{c4} + \omega_m + \mu) \left(I_{m(i)} + k_{3I_m}h\right), \\
k_{4I_c} &= \phi_c \left(E_{c(i)} + k_{3E_c}h\right) - (\epsilon\lambda_{m4} + \omega_c + \mu) \left(I_{c(i)} + k_{3I_c}h\right),
\end{aligned}$$

$$\begin{aligned}
k_{4I_m E_c} &= \delta \lambda_{c4} (I_m(i) + k_{3I_m} h) - (\sigma + \mu) (I_m E_c(i) + k_{3I_m E_c} h), \\
k_{4I_m c} &= \sigma (I_m E_c(i) + k_{3I_m E_c} h) + \phi_{mc} (E_{mc}(i) + k_{3E_{mc}} h) + \gamma (I_c E_m(i) + k_{3I_c E_m} h) - (\omega_{mc} + \mu) \\
&\quad (I_{mc}(i) + k_{3I_{mc}} h), \\
k_{4I_c E_m} &= \epsilon \lambda_{m4} (I_c(i) + k_{3I_c} h) - (\gamma + \mu) (I_c E_m(i) + k_{3I_c E_m} h), \\
k_{4S_v} &= \Lambda_v - (\lambda_{v4} + \mu_v) (S_v(i) + k_{3S_v} h), \\
k_{4E_v} &= \lambda_{v4} (S_v(i) + k_{3S_v} h) - (\phi_v + \mu_v) (E_v(i) + k_{3E_v} h), \\
k_{4I_v} &= \phi_v (E_v(i) + k_{3E_v} h) - (\mu_v) (I_v(i) + k_{3I_v} h).
\end{aligned}$$

RESULTS AND DISCUSSION

The numerical simulation results and the accuracy of the SEI models of the third-, and fourth-order Runge-Kutta methods are discussed in this section. Simulations are conducted using the MATLAB software and the calculation of the accuracy order using the Microsoft Excel software. Numerical simulations require initial values. Therefore, we use the third-order and fourth-order Runge-Kutta methods to solve equation (1) in the SEI model. The initial values of the variable and its parameters are listed in Table 3 and Table 4.

TABLE 3. Initial Values of Variables of the SEI model

Variables	Values
Human	
$N_{h(1)}$	$S_{h(1)} + E_{m(1)} + E_{c(1)} + E_{mc(1)} + I_{m(1)} + I_{c(1)} + I_{mE_c(1)} + I_{mc(1)} + I_{cE_m(1)}$
$S_{h(1)}$	2500
$E_{m(1)}$	1
$E_{c(1)}$	1
$E_{mc(1)}$	1
$I_{m(1)}$	10
$I_{c(1)}$	20
$I_{mE_c(1)}$	1
$I_{mc(1)}$	3
$I_{cE_m(1)}$	3
Mosquitoes (vector)	
$N_{v(1)}$	$S_{v(1)} + E_{v(1)} + I_{v(1)}$
$S_{v(1)}$	10000
$E_{v(1)}$	8
$I_{v(1)}$	10

The initial values of λ_c , λ_m , and λ_v are in the form of a predefined formula [36]. It means the three parameters have values that can change following the value of other related variables and the parameters. Wearing health masks, social and physical distance, hydro alcoholic gel, hand washing with soap, gathering intensity, and self-isolation were also included in this study as methods of community personal protection against COVID-19. This condition also brings up the efficacy of personal protection. A factor $(1 - \kappa\zeta)$ is used to represent the effect of personal protection, where $0 < \zeta < 1$ and $0 < \kappa < 1$. During the simulation, we selected initial values $\zeta = 0.5$ and $\kappa = 0.5$, which indicate that 50% of humans apply for personal protection. Then, the initial value of malaria transmission probability per mosquito bite, β_m selected 0.125 on numerical simulations.

TABLE 4. Initial Values of Parameters of the SEI model

Parameters	Values
Human	
Λ_h	10000
	$\frac{59 \times 365}{21}$
ϕ_m	0.8333
ϕ_c	0.6
ϕ_{mc}	0.333
ω_m	0.25
ω_c	0.3
ω_{mc}	0.025
μ	1
	$\frac{59 \times 365}{21}$
δ	1.02
ϵ	1.01
σ	0.4
γ	0.0833
κ	0 – 1
ζ	0 – 1
b	4.3×0.33
β_m	0.125 – 0.5
β_c	0.4531
λ_c	$\beta_c(1 - \kappa\zeta) \frac{(I_{c(1)} + I_{mc(1)} + I_{cE_m(1)})}{N_{h(1)}}$
λ_m	$\beta_m b \frac{I_{v(1)}}{N_{h(1)}}$
Mosquitoes	
Λ_v	$\frac{10^4}{21}$
ϕ_m	0.1
μ_v	$\frac{1}{21}$
β_v	0.48
λ_v	$\beta_v b \frac{(I_{m(1)} + I_{mc(1)} + I_{mE_c(1)})}{N_{h(1)}}$

Simulation Results of the Third-Order Runge-Kutta Method

The behavior of the co-dynamics of the spread of COVID-19 and malaria using the SEI model and the third-order Runge-Kutta method is depicted in Figure 1. The number of humans susceptible to malaria and COVID-19, namely, S_h , has decreased over time. That is because the number of individuals exposed to malaria only, namely E_m , increased at a certain period. That also happened to individuals exposed to COVID-19 only, namely E_c . However, at the same time, both subpopulations have decreased. The increasing number of individuals exposed to malaria or COVID-19 can open opportunities to influence the increase in the number of individuals infected. Initially, the number of individuals infected with malaria only, namely I_m , and individuals infected with COVID-19 only, namely I_c , decreased and stayed around a certain value at a certain period. Then the number increased over a long period and decreased again. Such behavior also occurs in the number of individuals infected with malaria and exposed to COVID-19, namely I_{mE_c} , and the number of individuals infected with COVID-19 and exposed to malaria, namely I_{cE_m} . These dynamics also occur in individuals exposed to malaria and COVID-19, namely E_{mc} , and individuals infected with malaria and COVID-19, namely I_{mc} , in the span of 0 to 60. Meanwhile, when close to infinity, $S_h(t)$, $E_m(t)$, $E_c(t)$, $E_{mc}(t)$, $I_m(t)$, $I_c(t)$, $I_{mE_c}(t)$, $I_{mc}(t)$, and $I_{cE_m}(t)$ will be constants. The human population will never become extinct.

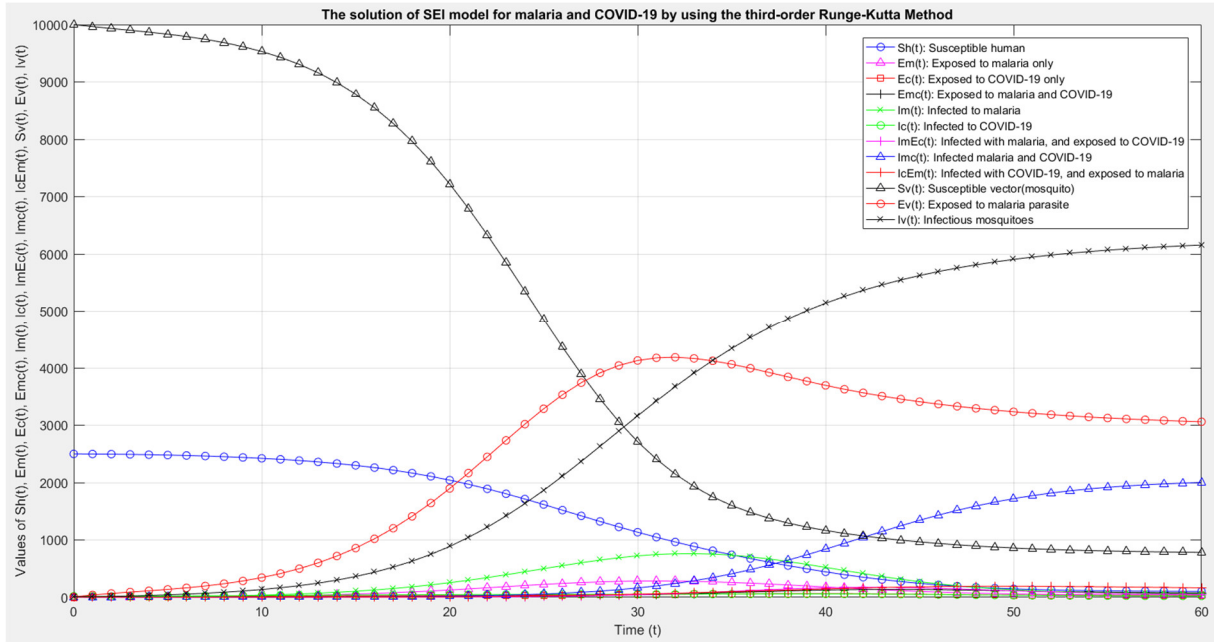


FIGURE 1. Graph of the solution to the spread of malaria and COVID-19 using the third-order Runge-Kutta method with $h = 1$ in the time range $[0, 60]$.

Furthermore, the number of mosquito populations (vectors) in each subpopulation has decreased or increased continuously over time. The number of susceptible mosquitoes, namely S_v , has decreased over time. Then the number of mosquitoes exposed to the malaria parasite, namely E_v , has increased over time. The number of mosquitoes that experienced an increase also occurred in the subpopulation of infectious mosquitoes, namely I_v . However, as time t approaches infinity, the number of mosquitoes will approach equilibrium. In other words, the mosquito population never extinct.

Simulation Results of the Fourth-Order Runge-Kutta Method

Figure 2 shows the results of the co-dynamics of the spread of malaria and COVID-19 with the SEI model using the fourth-order Runge-Kutta method. These results have similarities with the simulation results using the third-order Runge-Kutta method. The patterns of spread and behavior are also the same as those of the third-order Runge-Kutta method.

Method Accuracy Order

The solution patterns of the two methods are the same, as shown from the results of the simulations in Figure 1 and Figure 2. Meanwhile, one of the third-order and the fourth-order Runge-Kutta methods has better order of accuracy. Thus, it is necessary to measure the numerical error and the order of accuracy using the ODE45 algorithm in the MATLAB software for reference solution with ‘RelTol’ of 2.3×10^{-14} and ‘AbsTol’ of 1×10^{-15} . This is done because no exact analytical solution has been found for the SEI model in this paper.

To test numerical errors in each subpopulation, we use the absolute error formula. To calculate the order of accuracy for the time steps 1, 0.5, 0.25, 0.125, 0.0625, and 0.03125 we use the error results and the accuracy order formula is [42]

$$R_i = \frac{\log\left(\frac{E_i}{E_{i+1}}\right)}{\log\left(\frac{\Delta x_i}{\Delta x_{i+1}}\right)} \tag{11}$$

Here E_i is the error corresponding to the simulation using independent variable step Δx_i , and E_{i+1} is the error corresponding to the simulation using independent variable step Δx_{i+1} .

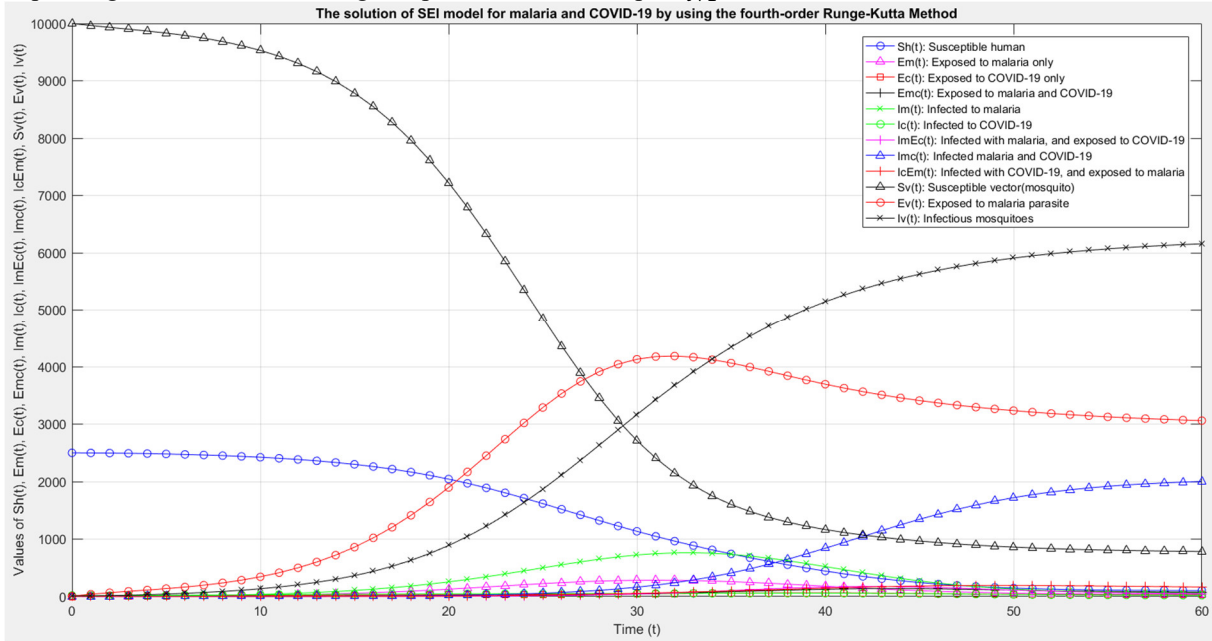


FIGURE 2. Graph of the solution to the spread of malaria and COVID-19 using the fourth-order Runge-Kutta method with $h = 1$ in the time range $[0, 60]$.

TABLE 5. Average error of the third-order Runge-Kutta method

h	S_h	E_m	E_c	E_{mc}	I_m	I_c	I_{mE_c}	I_{mc}	I_{cE_m}	S_v	E_v	I_v
1	14×10^{-2}	8×10^{-2}	4×10^{-2}	3×10^{-2}	15×10^{-2}	2×10^{-2}	5×10^{-2}	19×10^{-2}	3×10^{-2}	58×10^{-2}	48×10^{-2}	52×10^{-2}
0.5	20×10^{-3}	9×10^{-5}	4×10^{-3}	3×10^{-3}	19×10^{-3}	2×10^{-3}	6×10^{-3}	25×10^{-3}	3×10^{-3}	79×10^{-3}	62×10^{-3}	69×10^{-3}
0.25	26×10^{-4}	10×10^{-4}	4×10^{-4}	4×10^{-4}	24×10^{-4}	2×10^{-4}	7×10^{-4}	33×10^{-4}	4×10^{-4}	104×10^{-4}	79×10^{-4}	89×10^{-4}
0.125	34×10^{-5}	12×10^{-5}	5×10^{-5}	5×10^{-5}	30×10^{-5}	3×10^{-5}	9×10^{-5}	41×10^{-5}	5×10^{-5}	133×10^{-5}	100×10^{-5}	114×10^{-5}
0.0625	43×10^{-6}	15×10^{-6}	6×10^{-6}	7×10^{-6}	38×10^{-6}	3×10^{-6}	11×10^{-6}	52×10^{-6}	7×10^{-6}	168×10^{-6}	125×10^{-6}	143×10^{-6}
0.03125	54×10^{-7}	19×10^{-7}	7×10^{-7}	8×10^{-7}	48×10^{-7}	4×10^{-7}	14×10^{-7}	66×10^{-7}	9×10^{-7}	212×10^{-7}	157×10^{-7}	179×10^{-7}

TABLE 6. Average error of the fourth-order Runge-Kutta method

h	S_h	E_m	E_c	E_{mc}	I_m	I_c	I_{mE_c}	I_{mc}	I_{cE_m}	S_v	E_v	I_v
1	104×10^{-4}	186×10^{-4}	103×10^{-4}	31×10^{-4}	163×10^{-4}	77×10^{-4}	25×10^{-4}	49×10^{-4}	12×10^{-4}	463×10^{-4}	336×10^{-4}	135×10^{-4}
0.5	72×10^{-5}	93×10^{-5}	50×10^{-5}	13×10^{-5}	76×10^{-5}	34×10^{-5}	13×10^{-5}	34×10^{-5}	6×10^{-5}	280×10^{-5}	189×10^{-5}	98×10^{-5}
0.25	46×10^{-6}	51×10^{-6}	27×10^{-6}	6×10^{-6}	41×10^{-6}	18×10^{-6}	7×10^{-6}	22×10^{-6}	3×10^{-6}	173×10^{-6}	113×10^{-6}	66×10^{-6}

0.125	30×10^{-7}	30×10^{-7}	16×10^{-7}	3×10^{-7}	24×10^{-7}	10×10^{-7}	4×10^{-7}	14×10^{-7}	2×10^{-7}	108×10^{-7}	69×10^{-7}	42×10^{-7}
0.0625	19×10^{-8}	18×10^{-8}	10×10^{-8}	2×10^{-8}	14×10^{-8}	6×10^{-8}	3×10^{-8}	9×10^{-8}	1×10^{-8}	67×10^{-8}	43×10^{-8}	27×10^{-8}
0.03125	12×10^{-9}	11×10^{-9}	6×10^{-9}	1×10^{-9}	9×10^{-9}	4×10^{-9}	2×10^{-9}	6×10^{-9}	1×10^{-9}	42×10^{-9}	26×10^{-9}	17×10^{-9}

Table 5 and Table 6 show that the average error values in each subpopulation are different for each h . In addition, the average error values between subpopulations using the third-order Runge-Kutta method are larger than those using the fourth-order Runge-Kutta method.

TABLE 7. Accuracy Order of the third-order Runge-Kutta method

h	S_h	E_m	E_c	E_{mc}	I_m	I_c	I_{mE_c}	I_{mc}	I_{cE_m}	S_v	E_v	I_v
1	-	-	-	-	-	-	-	-	-	-	-	-
0.5	2.814	3.167	3.312	3.032	2.963	3.279	3.028	2.922	2.940	2.875	2.956	2.912
0.25	2.918	3.092	3.199	3.015	2.975	3.151	3.010	2.960	2.960	2.926	2.976	2.956
0.125	2.960	3.044	3.106	3.006	2.986	3.073	3.004	2.980	2.980	2.962	2.987	2.978
0.0625	2.980	3.021	3.054	3.003	2.993	3.035	3.002	2.990	2.990	2.981	2.993	2.989
0.03125	2.990	3.010	3.027	3.001	2.996	3.017	3.001	2.995	2.995	2.991	2.997	2.994

TABLE 8. Accuracy Order of the fourth-order Runge-Kutta method

h	S_h	E_m	E_c	E_{mc}	I_m	I_c	I_{mE_c}	I_{mc}	I_{cE_m}	S_v	E_v	I_v
1	-	-	-	-	-	-	-	-	-	-	-	-
0.5	3.856	4.320	4.367	4.596	4.413	4.486	4.280	3.849	4.365	4.047	4.153	3.788
0.25	3.945	4.167	4.190	4.342	4.215	4.256	4.126	3.925	4.152	4.015	4.067	3.903
0.125	3.975	4.086	4.098	4.189	4.111	4.135	4.061	3.963	4.071	4.005	4.032	3.953
0.0625	3.988	4.043	4.050	4.100	4.056	4.069	4.030	3.981	4.034	4.002	4.015	3.977
0.03125	3.994	4.022	4.025	4.052	4.028	4.035	4.015	3.991	4.017	4.001	4.008	3.989

Furthermore, Table 7 and Table 8 show that the accuracy of the third-order Runge-Kutta method is close to a certain value, namely 3, while the accuracy of the fourth-order Runge-Kutta method is close to 4. Then, the order accuracy results confirm that the fourth-order Runge-Kutta method's accuracy is better than the third-order Runge-Kutta method.

CONCLUSIONS

We have solved the SEI model effectively for the co-dynamic problem of the spread of malaria and COVID-19 using the third-, and the fourth-order Runge-Kutta methods. The solution from the simulation results of both methods using the MATLAB software show similarities. Therefore, it is necessary to look for the values of the numerical errors and the order of accuracy of each method to find the difference between the third-order and the fourth-order Runge-Kutta methods. The results obtained from our study confirmed that the fourth-order Runge-Kutta method is indeed more accurate than the third-order Runge-Kutta method in solving the co-dynamics of the spread of malaria and COVID-19 diseases with the SEI model.

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