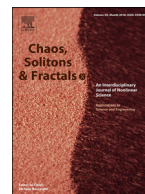




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## Mathematical analysis of a stochastic model for spread of Coronavirus

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### ABSTRACT

This paper is associated to investigate a stochastic SEIAQHR model for transmission of Coronavirus disease 2019 that is a recent great crisis in numerous societies. This stochastic pandemic model is established due to several safety protocols, for instance social-distancing, mask and quarantine. Three white noises are added to three of the main parameters of the system to represent the impact of randomness in the environment on the considered model. Also, the unique solvability of the presented stochastic model is proved. Moreover, a collocation approach based on the Legendre polynomials is presented to obtain the numerical solution of this system. Finally, some simulations are provided to survey the obtained results of this pandemic model and to identify the theoretical findings.

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### 1. Introduction

Using dynamical systems to describe behaviors of infectious diseases is an effective way to study how these diseases spread [1–3]. Mathematical models can greatly help researchers to better understand behaviors of deadly diseases around the world. The Susceptible-Infectious-Recovered (SIR) approach is a well-known basic model to analyze and to predict the epidemic of contagious diseases. Many researchers employed theoretical frameworks and numerical simulations to survey the manner of transmission of different infectious diseases [4–13].

In recent months, a specific group of viruses, named Coronavirus, has disrupted life around the world. Coronaviruses have a “crown” or corona of sugary-proteins. Thus, they have named Coronaviruses according to their specific appearance, in 1960. Special cases of Coronaviruses are the main sources of certain popular diseases such as the Middle East respiratory syndrome (MERS-CoV) and severe acute respiratory syndrome (SARS-CoV). Researches have shown that these viruses are conveyed between humans and animals. For example, MERS-CoV and SARS-CoV have transmitted from dromedary camels and civets to humans, respectively. Some

Coronavirus strains are only active in animals and have not yet infected human beings.

The Wuhan city in China was the first place that the Coronavirus disease 2019 (COVID-19) was identified. This new version of Coronaviruses was not previously diagnosed in human beings. Experts refer to different sources for this disease. Some of them believe the virus was first transmitted from pangolins or bats to humans in China. The high rate of contagion has caused that countries all over the world are affected by COVID-19. Cough, fever, difficulties in breathing and loss of sense of taste and smell are the main signs of the disease. High mortality rate and the serious economic and financial crisis created by the Coronavirus pandemic have motivated many researchers to study the behaviors of this virus and to find effective ways for facing this outbreak. Hence, some suitable susceptible-exposed-infectious-recovered (SEIR) type models are presented in these investigations to simulate the dynamic of the COVID-19 pandemic. The impact of asymptomatic category and quarantine on the transmission of COVID-19 [14], modeling the COVID-19 due to the interactivity between the animals, humans and the infections reservoir such as seafood market [15], investigating the dynamics of COVID-19 epidemics using real data from Pakistan [16], a model for the transmission of COVID-19 based on the Caputo-Fabrizio fractional order derivative [17], a stochastic model to study the stationary distribution and extinction of coronavirus epidemic [18], a stochastic model to evaluate health-care impact of Coronavirus in India [19], modeling the potential for mask use to curtail the COVID-19 [20], using fuzzy neural network

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models to predict COVID-19 time series in Mexico [21], investigation about the role of non-pharmaceutical interventions on reducing COVID-19 [22] and identifying the effect of social distancing in diminishing COVID-19 in Canada [23] are some examples of these scientific works.

In this work, first, we suggest a deterministic model to analyze the impact of preventive and health-protective measures such as quarantine, using masks and social distancing on the Coronavirus prevalence. A parameter called the force of infection is considered in the model. This parameter is affected by the factors related to the mentioned safety health protocols. Thus, this novel biological parameter helps us to verify the impact of safety strategies on the control of the disease. Many relationships between people are based on random contacts. So, by considering random parameters in the proposed deterministic model, a more realistic stochastic model for COVID-19 will be derived. Also, a new high-order numerical technique is introduced to find more accurate and reliable results for the presented biological system.

This paper has the following organization. In Section 2, a deterministic model for COVID-19 is presented based on some general strategies. Also, the analysis of this deterministic model will be developed. A stochastic model for the spread of Coronavirus is discussed in Section 3 and the existence of a unique positive solution for this system will be proved. In Section 4, a step-by-step collocation scheme is introduced and in Section 5, this method is employed to provide numerical simulations of the proposed stochastic model. At the end, the main conclusions are presented in Section 6.

## 2. The deterministic COVID-19 model

In the rest, the supposed population will be divided into several compartments to adapt a SEIR-type model describing the evolution of the COVID-19 pandemic. For simulation, the total size of the population under study is supposed to equal to  $N(t)$ .

In the survey conducted, every individual belongs to one of the following seven categories:

- ◆ the class S includes susceptible individuals who never infected before;
- ◆ the class E includes the individuals that have been newly-infected but those are not still sick and cannot transmit infection;
- ◆ the class I includes the infectious individuals with symptoms;
- ◆ the class A includes the infectious individuals that are not yet symptoms;
- ◆ the class Q includes the infectious individuals that are with slight symptoms and hence are at home quarantine;
- ◆ the class H includes the infectious individuals that are hospitalized;
- ◆ the class R includes the recovered individuals from the disease;

where

$$N(t) = S(t) + E(t) + I(t) + A(t) + Q(t) + H(t) + R(t).$$

$$\begin{aligned} \zeta_1 &:= c + \mu, & \zeta_2 &:= -c\rho(1 - q), & \zeta_3 &:= \delta_1 + \gamma_1 + \mu_1 + \mu, & \zeta_4 &:= -\alpha_1, \\ \zeta_5 &:= -\varphi\bar{\rho}, & \zeta_6 &:= -c(1 - \rho)(1 - q), & \zeta_7 &:= \gamma_A + \alpha_1 + \mu, & \zeta_8 &:= -\alpha_A, \\ \zeta_9 &:= -cq, & \zeta_{10} &:= \alpha_A + \varphi\bar{\rho} + \mu, & \zeta_{11} &:= -\delta_1, & \zeta_{12} &:= \mu + \gamma_H + \mu_1. \end{aligned}$$

Now, to introduce our model, we need to define some parameters. Let  $c$  and  $\theta$  show the contact and the transmission rates, respectively.  $q$  represents the rate of quarantine for exposed persons.  $\Lambda$  is the birth rate. To consider the effectiveness of social-distancing, the effective contact rate  $\beta$  is defined.  $\delta_1$  displays the transition rate from the symptomatic infected category to the quarantined infected category.  $\gamma_1$  and  $\gamma_A$  are the recovery rates of

symptomatic infected persons and asymptomatic infected persons, respectively and  $\gamma_H$  is the recovery rate of individuals in the class H. Also,  $\alpha_A$  and  $\alpha_1$  show the movement rate from the class Q to the class A and the transition rate from A to the class I, respectively.  $\rho$  and  $\mu_1$  represent the symptoms rate among infected individuals and the mortality rate due to Coronavirus, respectively. Moreover,  $c_m$  is the proportion of people who use masks and  $\varepsilon_m$  represents the effect of using mask to prevent catching the infection by susceptible individuals.  $0 \leq \theta_q \leq 1$  is the efficacy of quarantine and hospitalization admission in preventing virus transmission. Finally,  $\varphi$ ,  $\bar{\rho}$  and  $\mu$  represent the proportion of infected individuals with symptoms in quarantine, the movement rate from the category Q to I and the natural death rate, respectively. Let, the force of infection,  $\lambda$ , be defined as:

$$\lambda = \beta(1 - \varepsilon_m c_m) \frac{I + \theta A}{N - \theta_q(Q + H)}. \tag{1}$$

Then, according to the above variables and parameters, we have the following proposed system of equations for the dynamics of COVID-19:

$$\begin{cases} \dot{S} = \Lambda - (\lambda + \mu)S, \\ \dot{E} = \lambda S - (c + \mu)E, \\ \dot{I} = c\rho(1 - q)E + \varphi\bar{\rho}Q + \alpha_1 A - (\gamma_1 + \delta_1 + \mu_1 + \mu)I, \\ \dot{A} = c(1 - \rho)(1 - q)E + \alpha_A Q - (\alpha_1 + \gamma_A + \mu)A, \\ \dot{Q} = cqE - (\alpha_A + \varphi\bar{\rho} + \mu)Q, \\ \dot{H} = \delta_1 I - (\gamma_H + \mu_1 + \mu)H, \\ \dot{R} = \gamma_1 I + \gamma_A A + \gamma_H H - \mu R. \end{cases} \tag{2}$$

The diagram of this model is illustrated in Fig. 1.

The disease free equilibrium point of the model (2) can be obtained as

$$\mathcal{E}^0 = \left( \frac{\Lambda}{\mu}, 0, 0, 0, 0, 0, 0 \right)^T. \tag{3}$$

To assess the transmissibility of the disease, it is needed to get the basic reproduction number  $\mathcal{R}_0$ , i.e. the expected number of new infections that result from a single infectious person in a specific population of susceptible individuals. To determine this number for the COVID-19 model (2), we employ the method proposed in [24]. Suppose  $\varpi := \beta(1 - \varepsilon_m c_m)$ . The matrix of new infection terms  $\mathcal{F}$  and the matrix of the remaining transmission terms  $\mathcal{V}$  for the model (2) are defined as  $\mathcal{F}(\mathcal{E}^0) = [\mathbf{f}_{i,j}]_{5 \times 5}$  where

$$\mathbf{f}_{i,j} = \begin{cases} \varpi, & i = 1, j = 2, \\ \theta\varpi, & i = 1, j = 3, \\ 0, & o.w., \end{cases}$$

and

$$\mathcal{V}(\mathcal{E}^0) = \begin{pmatrix} \zeta_1 & 0 & 0 & 0 & 0 \\ \zeta_2 & \zeta_3 & \zeta_4 & \zeta_5 & 0 \\ \zeta_6 & 0 & \zeta_7 & \zeta_8 & 0 \\ \zeta_9 & 0 & 0 & \zeta_{10} & 0 \\ 0 & \zeta_{11} & 0 & 0 & \zeta_{12} \end{pmatrix},$$

in which

The required  $\mathcal{R}_0$  is the spectral radius of the matrix  $\mathcal{F}\mathcal{V}^{-1}$ . Hence, we get

$$\mathcal{R}_0 = \frac{\varpi c}{\gamma \eta \xi (c + \mu)} \left( \xi \bar{\rho} + \eta \theta \tilde{\beta} + q \bar{\delta} \right), \tag{4}$$

where  $\tilde{\theta} = (1 - \rho)(1 - q)$  and

$$\gamma = \alpha_1 + \gamma_A + \mu, \quad \eta = \delta_1 + \gamma_1 + \mu_1 + \mu, \quad \xi = \varphi\bar{\rho} + \alpha_A + \mu,$$

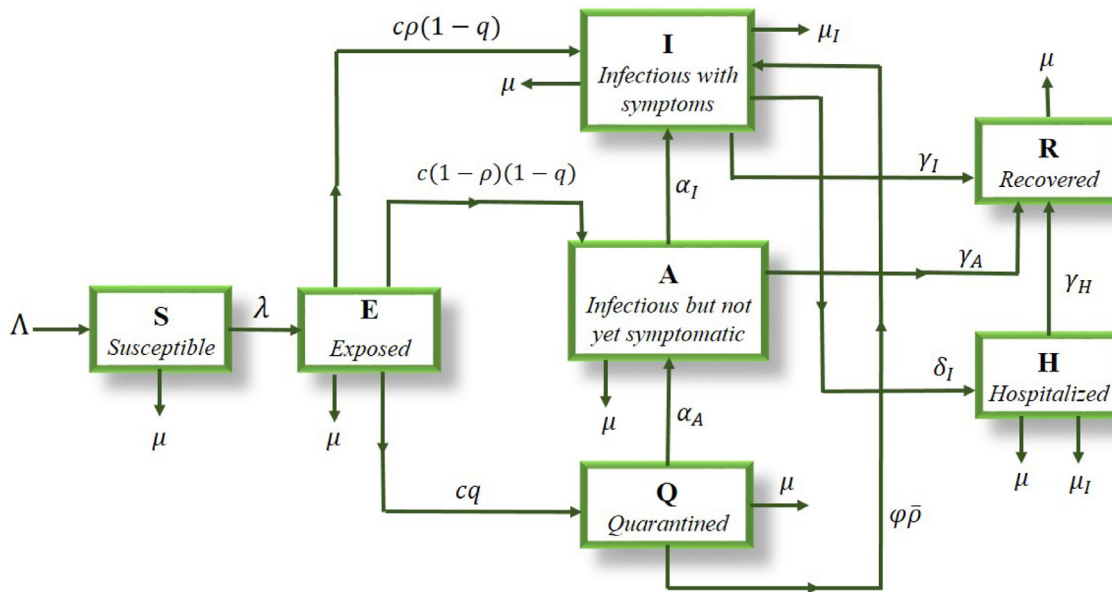


Fig. 1. The diagram of the proposed coronavirus transmission model.

$\bar{q} = \gamma \rho(1 - q) + \tilde{\theta} \alpha_1$ ,  $\tilde{\beta} = \xi \tilde{\theta} + q \alpha_A$ ,  $\tilde{\delta} = \alpha_A \alpha_1 + \gamma \varphi \bar{\rho}$ . Also, the endemic equilibrium point  $\mathcal{E}_* = (S^*, E^*, I^*, A^*, Q^*, H^*, R^*)^T$  is determined by solving the system

$$\begin{cases} \Lambda - (\lambda + \mu)S^* = 0, \\ \lambda S^* - (c + \mu)E^* = 0, \\ c \dot{\theta} E^* + \varphi \bar{\rho} Q^* + \alpha_1 A^* - \eta I^* = 0, \\ c \dot{\alpha} E^* + \alpha_A Q^* - \gamma A^* = 0, \\ c q E^* - \xi Q^* = 0, \\ \delta_1 I^* - \zeta H^* = 0, \\ \gamma_1 I^* + \gamma_A A^* + \gamma_H H^* - \mu R^* = 0, \end{cases} \quad (5)$$

where  $\zeta = \mu_1 + \gamma_H + \mu$ ,  $\dot{\theta} = \rho(1 - q)$  and  $\dot{\alpha} = (1 - \rho)(1 - q)$ . So

$$\begin{aligned} S^* &= \Psi_S E^*, & I^* &= \Psi_I E^*, \\ Q^* &= \Psi_Q E^*, & H^* &= \Psi_H E^*, \\ A^* &= \Psi_A E^*, & R^* &= \Psi_R E^*, \end{aligned}$$

where

$$E^* = \frac{\Lambda(\Lambda \mathcal{R}_0 - \mu)}{(c + \mu)(\Lambda \mathcal{R}_0 - \mu) + (1 - \theta_q)(\Psi_Q + \Psi_H) + \Psi_I + \Psi_R + \mu^2 + 1},$$

and

$$\begin{aligned} \Psi_S &= \frac{c + \mu}{\lambda}, & \Psi_Q &= \frac{c q}{\xi}, \\ \Psi_A &= \frac{c}{\xi \gamma} (\dot{\alpha} \xi + \alpha_A q), & \Psi_I &= \frac{c}{\eta \xi \gamma} (\dot{\theta} \xi \gamma + \varphi \bar{\rho} q \gamma + \alpha_1 \xi \dot{\alpha} + \alpha_A \alpha_1 q), \\ \Psi_H &= \frac{\delta_1}{\omega} \Psi_I, & \Psi_R &= \frac{1}{\mu} \left( \gamma_A \Psi_A + \left( \gamma_1 + \frac{\gamma_H \delta_1}{\zeta} \right) \Psi_I \right). \end{aligned}$$

### 3. Stochastic model

Real-world models are usually influenced by uncertain natural factors. Thus, incorporating environmental interactions helps to find a better picture of the COVID-19 pandemic in a society. So, it is more appropriate to present the model of disease by stochastic differential equations. In the rest, we improve the model (2) by considering some noisy environmental effects. To this end, the Brownian motion process will be employed to simulate randomness in data.

In the spread of disease model (2), we focus on the three transmission rates  $\beta$ ,  $\alpha_I$  and  $\gamma_H$ . In the environment, these parameters

are not fixed but oscillate around some average values. Let some white noises be added to them as

$$\begin{aligned} \beta &\rightarrow \beta + \sigma_1 \dot{B}_1(t), \\ \alpha_1 &\rightarrow \alpha_1 + \sigma_2 \dot{B}_2(t), \\ \gamma_H &\rightarrow \gamma_H + \sigma_3 \dot{B}_3(t), \end{aligned}$$

where  $B_i(t)$ ,  $i = 1, 2, 3$ , are standard Brownian motions [25,26] and  $\sigma_i$ ,  $i = 1, 2, 3$ , are the intensities of environmental oscillations that  $\sigma_i^2 > 0$ . Now, we obtain the following stochastic model as

$$\begin{cases} dS = (\Lambda - (\lambda + \mu)S)dt - \frac{\sigma_1}{\beta} \lambda S dB_1(t), \\ dE = (\lambda S - (c + \mu)E)dt + \frac{\sigma_1}{\beta} \lambda S dB_1(t), \\ dI = (c\rho(1 - q)E + \varphi \bar{\rho} Q + \alpha_1 A - (\gamma_1 + \delta_1 + \mu_1 + \mu)I)dt \\ \quad + \sigma_2 A dB_2(t), \\ dA = (c(1 - \rho)(1 - q)E + \alpha_A Q - (\alpha_1 + \gamma_A + \mu)A)dt \\ \quad - \sigma_2 A dB_2(t), \\ dQ = (cqE - (\alpha_A + \varphi \bar{\rho} + \mu)Q)dt, \\ dH = (\delta_1 I - (\gamma_H + \mu_1 + \mu)H)dt - \sigma_3 H dB_3(t), \\ dR = (\gamma_1 I + \gamma_A A + \gamma_H H - \mu R)dt + \sigma_3 H dB_3(t). \end{cases} \quad (6)$$

To investigate this improved model, first, we prove the existence of unique global positive solution for this system. Let

$$\mathbb{R}_+^7 := \{(u_1, \dots, u_7) \in \mathbb{R}^7 \mid u_i > 0, \quad i = 1, \dots, 7\}.$$

**Theorem 1.** For any initial value  $(S(0), E(0), I(0), A(0), Q(0), H(0), R(0)) \in \mathbb{R}_+^7$  and every  $t \geq 0$ , the system (6) have an unique solution. This solution will remain positive with probability one.

**Proof.** The coefficients of system (6) are locally Lipschitz continuous. Hence, this system has a unique local solution

$$(S(t), E(t), I(t), A(t), Q(t), H(t), R(t)),$$

on  $[0, \tau_e)$ , where  $\tau_e$  is the “explosion time” [25], for any initial value

$$(S(0), E(0), I(0), A(0), Q(0), H(0), R(0)) \in \mathbb{R}_+^7.$$

Now, we show that  $\tau_e = \infty$  almost surely (a.s.) and as a result the solution is global. Suppose  $\omega_0 > 0$  is sufficiently large such that the initial values  $S(0), E(0), I(0), A(0), Q(0), H(0)$  and  $R(0)$  lie within the interval  $[\frac{1}{\omega_0}, \omega_0]$ . Let

$$\begin{aligned} \rho_{\min}(t) &:= \min\{S(t), E(t), I(t), A(t), Q(t), H(t), R(t)\}, \\ \rho_{\max}(t) &:= \max\{S(t), E(t), I(t), A(t), Q(t), H(t), R(t)\}, \end{aligned}$$

for any  $w \geq \omega_0$ , and

$$t_w = \inf \left\{ t \in [0, t_e) : \rho_{\min}(t) \leq \frac{1}{w} \text{ or } \rho_{\max}(t) \geq w \right\}. \quad (7)$$

Set  $\inf \emptyset = \infty$  where  $\emptyset$  shows the empty set. From this definition it can be concluded that  $t_w$  is increasing when  $w \rightarrow \infty$ . Let  $t_\infty = \lim_{w \rightarrow \infty} t_w$ . So,  $t_\infty \leq t_0$  a.s. Therefore, if we can prove  $t_\infty = \infty$  a.s., then  $t_e = \infty$  a.s. and

$$(S(t), E(t), I(t), A(t), Q(t), H(t), R(t)) \in \mathbb{R}_+^7 \text{ a.s., } \forall t \geq 0.$$

To complete the proof, it is sufficient to prove that  $t_\infty = \infty$  a.s. Let this is not correct. Therefore, there exists two constants  $\hat{\rho} > 0$  and  $\tilde{\kappa} \in (0, 1)$  such that  $\mathbb{P}\{t_\infty \leq \hat{\rho}\} > \tilde{\kappa}$ . So

$$\exists \omega_1 \in \mathbb{Z}, \omega_1 > \omega_0, \text{ s.t. } \mathbb{P}\{t_w \leq \hat{\rho}\} \geq \tilde{\kappa}, \quad \forall w \geq \omega_1. \quad (8)$$

For  $t \leq t_w$

$$dN(t) = \left( \Lambda - \mu N(t) - \mu_1(I(t) + H(t)) \right) dt \leq (\Lambda - \mu N(t)) dt.$$

Thus

$$N(t) = \begin{cases} \frac{\Lambda}{\mu}, & \text{if } N(0) \leq \frac{\Lambda}{\mu}, \\ N(0), & \text{if } N(0) > \frac{\Lambda}{\mu}, \end{cases} := \tilde{J}.$$

Consider the twice differentiable function  $\psi : \mathbb{R}_+^7 \rightarrow \mathbb{R}_+$  with the definition

$$\begin{aligned} \psi(S, E, I, A, Q, H, R) := & (S - 1 - \log S) + (E - 1 - \log E) \\ & + (I - 1 - \log I) \\ & + (A - 1 - \log A) + (Q - 1 - \log Q) \\ & + (H - 1 - \log H) + (R - 1 - \log R). \end{aligned}$$

Since  $\log u \leq u - 1$ , for every  $u \geq 0$ , hence,  $\psi$  is nonnegative. Due to the system (6) and Itô's formula on  $\psi$

$$\begin{aligned} d\psi(S, E, I, A, Q, H, R) = & \mathcal{L}\psi(S, E, I, A, Q, H, R)dt \\ & + \sigma_1 \frac{\lambda}{\beta} \left(1 - \frac{S}{E}\right) dB_1(t) \\ & + \sigma_2 \left(1 - \frac{A}{I}\right) dB_2(t) + \sigma_3 \left(1 - \frac{H}{R}\right) dB_3(t), \end{aligned}$$

where

$$\begin{aligned} \mathcal{L}\psi(S, E, I, A, Q, H, R) = & \left(1 - \frac{1}{S}\right) (\Lambda - (\lambda + \mu)S) + \frac{1}{2\beta^2} \sigma_1^2 \lambda^2 \\ & + \left(1 - \frac{1}{E}\right) (\lambda S - (c + \mu)E) + \frac{1}{2\beta^2} \sigma_1^2 \lambda^2 \frac{S^2}{E^2} + \frac{1}{2} \sigma_2^2 \frac{A^2}{I^2} \\ & + \left(1 - \frac{1}{I}\right) (c\rho(1 - q)E + \varphi\bar{\rho}Q + \alpha_1 A - (\gamma_1 + \delta_1 + \mu_1 + \mu)I) \\ & + \frac{1}{2} \sigma_2^2 \\ & + \left(1 - \frac{1}{A}\right) (c(1 - \rho)(1 - q)E + \alpha_A Q - (\alpha_1 + \gamma_A + \mu)A) \\ & + \left(1 - \frac{1}{Q}\right) (cqE - (\alpha_A + \varphi\bar{\rho} + \mu)Q) + \left(1 - \frac{1}{H}\right) \\ & \times (\delta_1 I - (\gamma_H + \mu_1 + \mu)H) \\ & + \frac{1}{2} \sigma_3^2 + \left(1 - \frac{1}{R}\right) (\gamma_1 I + \gamma_A A + \gamma_H H - \mu R) + \frac{1}{2} \sigma_3^2 \frac{H^2}{R^2} \\ = & \Lambda + \lambda + c + \gamma_1 + \delta_1 + \mu_1 + \alpha_1 + \gamma_A + \alpha_A + \varphi\bar{\rho} + \gamma_H + \mu_1 + 7\mu \\ & + \frac{1}{2\beta^2} \sigma_1^2 \lambda^2 \left(1 + \frac{S^2}{E^2}\right) + \frac{\sigma_2^2}{2} \left(1 + \frac{A^2}{I^2}\right) + \frac{\sigma_3^2}{2} \left(1 + \frac{H^2}{R^2}\right) - \mu N - \mu_1(I + H) \\ & - \frac{\Lambda}{S} - c\rho(1 - q)\frac{E}{I} - \varphi\bar{\rho}\frac{Q}{I} - \alpha_1\frac{A}{I} - c(1 - \rho)(1 - q)\frac{E}{A} \\ & - \alpha_A\frac{Q}{A} - cq\frac{E}{Q} - \delta_1\frac{1}{H} - \gamma_1\frac{1}{R} - \gamma_A\frac{A}{R} - \gamma_H\frac{H}{R} \\ \leq & \Lambda + \lambda + c + \gamma_1 + \delta_1 + \mu_1 + \alpha_1 + \gamma_A + \alpha_A + \varphi\bar{\rho} + \gamma_H + \mu_1 \\ & + 7\mu + \frac{1}{2\beta^2} \sigma_1^2 \lambda^2 (S^2 + E^2) + \frac{\sigma_2^2}{2} (A^2 + I^2) + \frac{\sigma_3^2}{2} (H^2 + R^2) \\ \leq & \Lambda + \lambda + c + \gamma_1 + \delta_1 + \mu_1 + \alpha_1 + \gamma_A + \alpha_A + \varphi\bar{\rho} + \gamma_H \\ & + \mu_1 + 7\mu + \left(\frac{1}{\beta^2} \sigma_1^2 \lambda^2 + \sigma_2^2 + \sigma_3^2\right) \tilde{J}^2 := \mathcal{Y}, \end{aligned}$$

which is bounded and  $\mathcal{Y} \in \mathbb{R}_+$ . Therefore, we obtain

$$\begin{aligned} \int_0^{t_w \wedge \hat{\rho}} d\psi(S(t), E(t), I(t), A(t), Q(t), H(t), R(t)) \leq & \\ \int_0^{t_w \wedge \hat{\rho}} \mathcal{Y} dt + \int_0^{t_w \wedge \hat{\rho}} \sigma_1 \frac{\lambda}{\beta} \left(1 - \frac{S}{E}\right) dB_1(t) & \\ + \int_0^{t_w \wedge \hat{\rho}} \sigma_2 \left(1 - \frac{A}{I}\right) dB_2(t) + \int_0^{t_w \wedge \hat{\rho}} \sigma_3 \left(1 - \frac{H}{R}\right) dB_3(t), & \end{aligned}$$

and

$$\begin{aligned} & \mathbb{E}(\psi(S(t), E(t), I(t), A(t), Q(t), H(t), R(t))) \\ & \leq \mathbb{E}(\psi(S(0), E(0), I(0), A(0), Q(0), H(0), R(0))) + \mathcal{Y}\mathbb{E}(t) \\ & \leq \mathbb{E}(\psi(S(0), E(0), I(0), A(0), Q(0), H(0), R(0))) + \mathcal{Y}\hat{\rho}, \quad (9) \end{aligned}$$

where  $(\cdot) = (t_w \wedge \hat{\rho})$  and  $\mathbb{E}$  shows the mathematical expectation. Let

$$\Omega_w := \{t_w \leq \hat{\rho}\}, \quad w \geq \omega_1.$$

From Eq. (8), we have  $\mathbb{P}(\Omega_w) \geq \tilde{\kappa}$ . Moreover, for every  $v \in \Omega_w$  at least one of the variables  $S, E, I, A, Q, H$  or  $R$  is greater or equal to  $w$ , or less than or equal to  $\frac{1}{w}$ . Hence,  $\Pi_{t_w} := \psi(S(t_w), E(t_w), I(t_w), A(t_w), Q(t_w), H(t_w), R(t_w))$  is not less than  $w - 1 - \log w$  or  $\frac{1}{w} - 1 + \log w$ , i.e.,

$$\Pi_{t_w} \geq \left(w - 1 - \log w\right) \wedge \left(\frac{1}{w} - 1 + \log w\right).$$

Then, (8) and (9) results

$$\begin{aligned} \mathbb{E}(\Pi_0) + \mathcal{Y}\hat{\rho} \geq & \mathbb{E}(\mathcal{I}_{\Omega_w} \Pi_{t_w}) \\ \geq & \tilde{\kappa} \left[ \left(w - 1 - \log w\right) \wedge \left(\frac{1}{w} - 1 + \log w\right) \right], \end{aligned}$$

where  $\mathcal{I}_{\Omega_w}$  is the usual notation for indicator of set  $\Omega_w$ . If  $w \rightarrow \infty$ , then

$$\infty > \mathbb{E}(\Pi_0) + \mathcal{Y}\hat{\rho} = \infty,$$

which is a contradiction. So, the hypothesis  $\mathbb{P}\{t_\infty \leq \hat{\rho}\} > \tilde{\kappa}$  is wrong and  $t_\infty = \infty$  a.s.  $\square$

### 3.1. Exponentially stability

In this subsection, we study the stability of disease-free equilibrium for model (6). Let  $\mathcal{C}^{2,1}(\mathbb{R}^n \times \mathbb{R})$  is the family of non-negative functions  $\psi(u, t)$  on  $\mathbb{R}^n \times \mathbb{R}$  that are twice continuously differentiable in  $u$  and once in  $t$ .

**Theorem 2.** [27] Suppose there exists a function  $\psi(u, t) \in \mathcal{C}^{2,1}(\mathbb{R}^n \times \mathbb{R})$  satisfying the following inequalities

$$|\psi(u, t)| \leq \zeta_1 |u|^p, \quad \mathcal{L}\psi(u, t) \leq -\zeta_2 |u|^p,$$

where  $\zeta_i > 0, i = 1, 2$  and  $p > 0$ . Thus, the equilibrium point  $\mathcal{E}^0$  of the system (6) is exponentially  $p$ -stable. Also,  $\mathcal{E}^0$  is exponentially stable in mean square and globally asymptotically stable, when  $p = 2$ .

**Lemma 1.** [4] Suppose  $p \geq 2$  and  $z, v \in \mathbb{R}_+$  and  $\varepsilon > 0$ . Thus, we have

$$\begin{aligned} z v^{p-1} & \leq \frac{\varepsilon^{1-p}}{p} z^p + \frac{(p-1)\varepsilon}{p} v^p, \\ z^2 v^{p-2} & \leq \frac{2\varepsilon^{\frac{2-p}{2}}}{p} z^p + \frac{(p-2)\varepsilon}{p} v^p. \end{aligned}$$

**Theorem 3.** Let  $p \geq 2$ , the disease-free equilibrium point

$$\mathcal{E}^0 = \left(\frac{\Lambda}{\mu}, 0, 0, 0, 0, 0, 0\right) \in \mathbb{R}_+^7$$

of the system (6) is exponentially  $p$ -stable, if

$$\begin{aligned} \frac{\sigma_2^2}{2} (p-1) & < \alpha_1 + \gamma_A + \mu, \\ \text{and } \frac{\sigma_3^2}{2} (p-1) & < \gamma_H + \mu_1 + \mu. \end{aligned}$$

**Proof.** Considering the following Lyapunov function

$$\psi = \frac{1}{p} \left( \left(\frac{\Lambda}{\mu} - S\right)^p + E^p + I^p + A^p + Q^p + H^p + R^p \right),$$

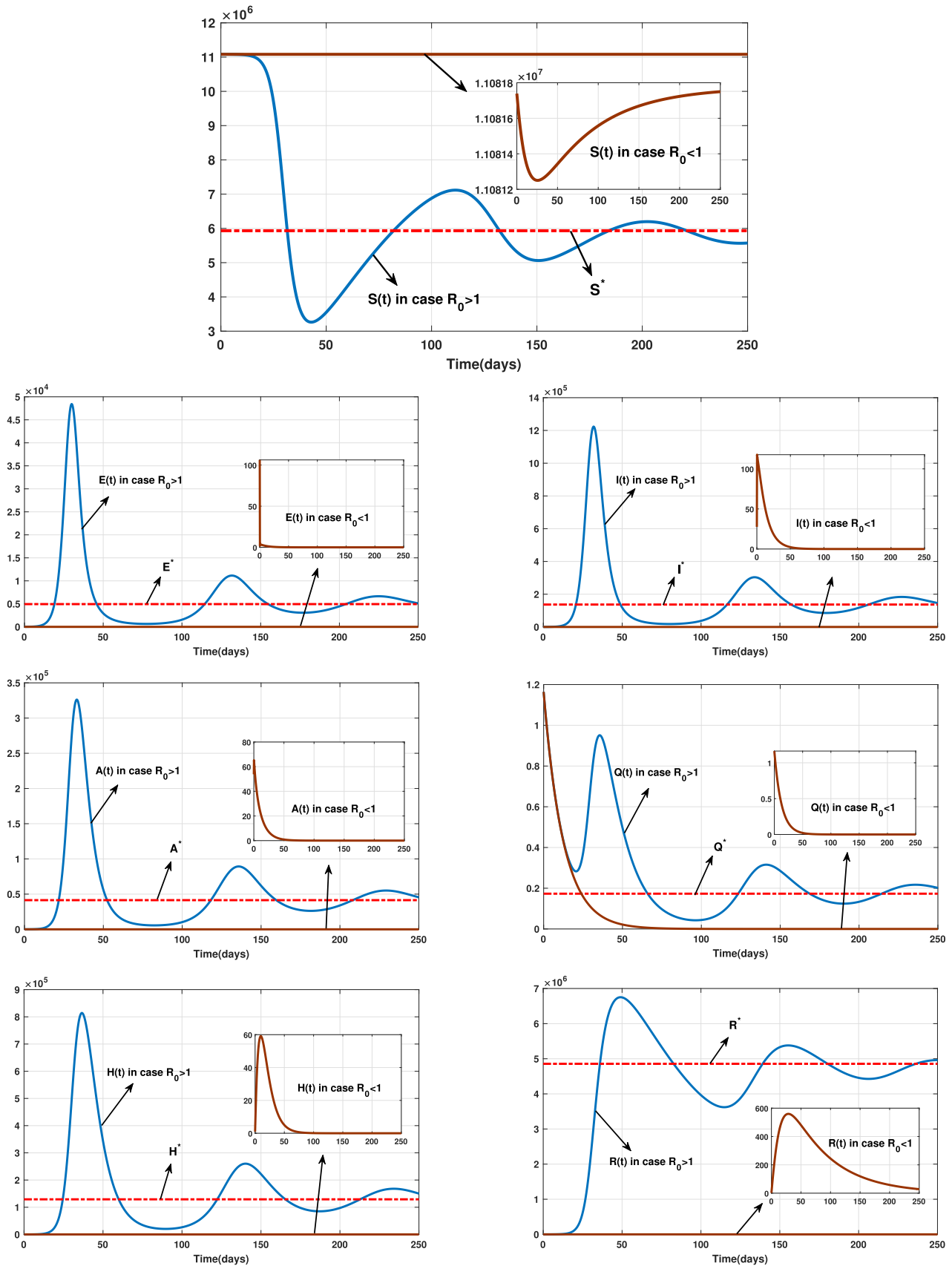


Fig. 2. The trajectories of the solution for the cases  $R_0 < 1$  and  $R_0 > 1$ , when  $\sigma_1 = \sigma_2 = \sigma_3 = 0$ .

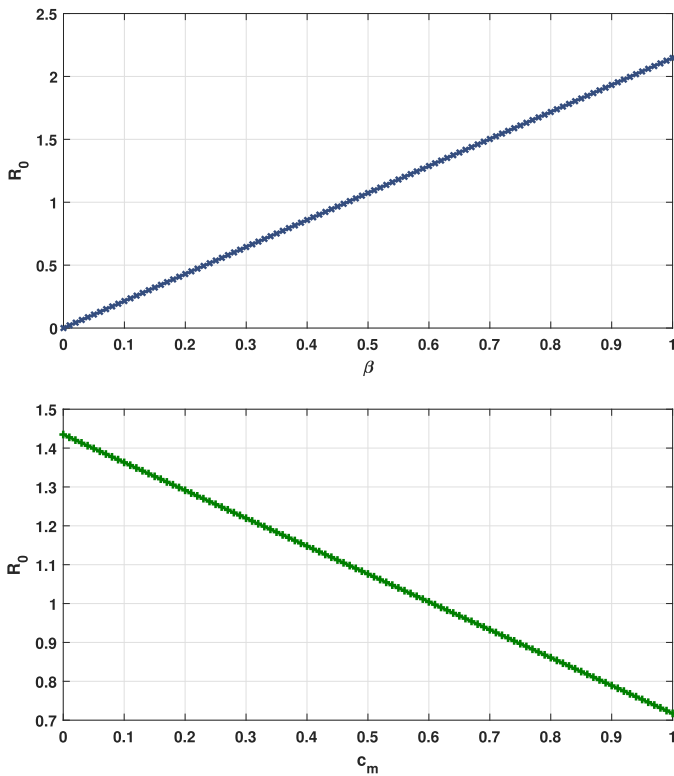


Fig. 3. Reproduction number  $\mathcal{R}_0$  for several values of  $\beta$  (up) and  $c_m$  (down).

where  $p \geq 2$ , then

$$\begin{aligned} \mathcal{L}\psi = & -\left[\frac{\Lambda\mu}{\Lambda-\mu S}\left(\frac{\Lambda}{\mu}-S\right)^p + (c+\mu)E^p + (\gamma_1 + \delta_1 + \mu_1 + \mu)I^p\right. \\ & + (\alpha_1 + \gamma_A + \mu)A^p + (\alpha_A + \varphi\bar{\rho} + \mu)Q^p + (\gamma_H + \mu_1 + \mu)H^p + \mu R^p \\ & + (\lambda + \mu)S\left(\frac{\Lambda}{\mu}-S\right)^{p-1} + \lambda SE^{p-1} + c\rho(1-q)EI^{p-1} + \varphi\bar{\rho} QI^{p-1} + \alpha_1 AI^{p-1} \\ & + c(1-\rho)(1-q)EA^{p-1} + \alpha_A QA^{p-1} + cQE^{p-1} + \delta_1 HI^{p-1} + \gamma_1 IR^{p-1} \\ & + \gamma_A AR^{p-1} + \gamma_H HR^{p-1} + \frac{\sigma_1^2}{2\beta^2}\lambda^2(p-1)S^2\left(\frac{\Lambda}{\mu}-S\right)^{p-2} + \frac{\sigma_1^2}{2\beta^2}\lambda^2(p-1)S^2E^{p-2} \\ & \left. + \frac{\sigma_2^2}{2}(p-1)A^2I^{p-2} + \frac{\sigma_2^2}{2}(p-1)A^p + \frac{\sigma_2^2}{2}(p-1)H^p + \frac{\sigma_2^2}{2}(p-1)H^2R^{p-2}\right. \end{aligned}$$

By using Lemma 1, we have

$$\begin{aligned} \mathcal{L}\psi \leq & -\left[\frac{\Lambda\mu}{\Lambda-\mu S}\left(\frac{\Lambda}{\mu}-S\right)^p + (c+\mu)E^p + (\gamma_1 + \delta_1 + \mu_1 + \mu)I^p\right. \\ & + (\alpha_1 + \gamma_A + \mu)A^p + (\alpha_A + \varphi\bar{\rho} + \mu)Q^p + (\gamma_H + \mu_1 + \mu)H^p + \mu R^p \\ & + \left(\lambda + \mu\right)\frac{(p-1)\varepsilon}{p} + \frac{\sigma_1^2}{2\beta^2}\lambda^2(p-1)(p-2)\varepsilon\left(\frac{\Lambda}{\mu}-S\right)^p \\ & + \left(2\lambda + \mu\right)\frac{\varepsilon^{1-p}}{p} + \frac{2\sigma_1^2}{\beta^2}\lambda^2(p-1)\varepsilon^{\frac{2-p}{2}}\left)S^p\right. \\ & + \left(\frac{(p-1)}{p}\left(\lambda + \frac{\sigma_1^2}{2\beta^2}\lambda^2(p-2)\varepsilon + \frac{c}{p}\varepsilon^{1-p}\right)E^p\right. \\ & + \left(\frac{(p-1)}{p}\left(c\rho(1-q) + \varphi\bar{\rho} + \alpha_1 + \frac{\sigma_2^2}{2}(p-2)\varepsilon + (\delta_1 + \gamma_1)\frac{\varepsilon^{1-p}}{p}\right)I^p\right. \\ & + \left(\alpha_1 + \gamma_A\right)\frac{\varepsilon^{1-p}}{p} + (\alpha_A + c(1-\rho)(1-q))\frac{(p-1)\varepsilon}{p} + \frac{\sigma_2^2}{p}(p-1)\varepsilon^{\frac{2-p}{2}} \\ & + \frac{\sigma_2^2}{2}(p-1)A^p + (cq)\frac{(p-1)\varepsilon}{p} + (\varphi\bar{\rho} + \alpha_A)\frac{\varepsilon^{1-p}}{p}Q^p \\ & + \left(\delta_1\frac{(p-1)\varepsilon}{p} + \gamma_H\frac{\varepsilon^{1-p}}{p} + \sigma_3^2(p-1)\frac{\varepsilon^{\frac{2-p}{2}}}{p} + \frac{\sigma_2^2}{2}(p-1)\right)H^p \\ & \left. + \left((\gamma_1 + \gamma_A + \gamma_H)\frac{(p-1)\varepsilon}{p} + \frac{\sigma_3^2}{2\beta}(p-1)(p-2)\varepsilon\right)R^p\right. \end{aligned}$$

According to the assumption, we know  $\frac{\sigma_2^2}{2}(p-1) < \alpha_1 + \gamma_A + \mu$  and  $\frac{\sigma_2^2}{2}(p-1) < \gamma_H + \mu_1 + \mu$ . Let  $\varepsilon$  is sufficiently small as the coefficients  $(\frac{\Lambda}{\mu}-S)^p$ ,  $E^p$ ,  $I^p$ ,  $A^p$ ,  $Q^p$ ,  $H^p$  and  $R^p$  be negative. Thus, from Theorem 2,  $\mathcal{E}^0$  is exponentially p-stable. If the conditions  $\frac{\sigma_2^2}{2} < \alpha_1 + \gamma_A + \mu$  and  $\frac{\sigma_3^2}{2} < \gamma_H + \mu_1 + \mu$  hold, then, in the special case  $p = 2$ , the equilibrium point  $\mathcal{E}^0$  of system (6) will be globally asymptotically stable.  $\square$

### 4. Numerical Scheme

In this section, we describe a step-by-step collocation approach based on the Legendre polynomials to solve a system of nonlinear stochastic differential equations.

**Definition 1.** [28] The recurrence formula of Legendre polynomials on interval  $[-1, 1]$  are defined as

$$\theta_{i+1}(t) = \frac{1}{i+1}\{(2i+1)t\theta_i(t) - i\theta_{i-1}(t)\}, \quad i = 1, 2, \dots,$$

where  $\theta_0(t) = 1$  and  $\theta_1(t) = t$ .

**Definition 2.** The shifted Legendre polynomials on  $[t_{\min}, t_{\max}]$  are defined as

$$t_{\min}^{\tau} t_{\max}^{\theta} \theta_i = \theta_i\left(\frac{2}{t_{\max}-t_{\min}}(t-t_{\min})+1\right), \quad i = 0, 1, 2, \dots$$

We use a step-by-step approach to solve the system (6) on the interval  $[0, T_{\max}]$ . For this purpose, let  $\varepsilon > 0$  and  $\bar{P} := [\frac{T_{\max}}{\varepsilon}]$ . Now, to find a numerical solution of the system on the subinterval  $[0, \varepsilon]$ , we will employ a Legendre collocation method. Consider the numerical solutions  ${}_1\mathcal{X}_n^i(t)$ ,  $i = 1, \dots, 7$ , as follows

$${}_1\mathcal{X}_n^i(t) = \sum_{j=0}^n {}_0^e x_j^i \theta_j(t) = {}_0^e \mathbf{X}_i^T \theta(t), \quad i = 1, \dots, 7, \tag{10}$$

where  $S(t) \simeq {}_1\mathcal{X}_n^1(t)$ ,  $E(t) \simeq {}_1\mathcal{X}_n^2(t)$ ,  $I(t) \simeq {}_1\mathcal{X}_n^3(t)$ ,  $A(t) \simeq {}_1\mathcal{X}_n^4(t)$ ,  $Q(t) \simeq {}_1\mathcal{X}_n^5(t)$ ,  $H(t) \simeq {}_1\mathcal{X}_n^6(t)$ ,  $R(t) \simeq {}_1\mathcal{X}_n^7(t)$  and

$$\begin{aligned} \mathbf{X}_i^0 & := [x_{00}^i, \dots, x_{0j}^i, \dots, x_{0n}^i]^T, \\ \Theta_0^e(t) & := [\phi_0^e(t), \dots, \phi_j^e(t), \dots, \phi_n^e(t)]^T, \end{aligned}$$

for  $t \in [0, \varepsilon]$ . According to (6) and (10), we have

$$\begin{cases} \Psi_1^1(t) \triangleq {}_0^e \mathbf{X}_1^T \theta(t) - \Lambda + ({}_0^e \Pi(t) + \mu) {}_0^e \mathbf{X}_1^T \theta(t) \\ \quad + \frac{\sigma_1}{\beta} {}_0^e \Pi(t) {}_0^e \mathbf{X}_1^T \theta(t) \mathcal{B}_1(t) \simeq 0, \\ \Psi_2^1(t) \triangleq {}_0^e \mathbf{X}_2^T \theta(t) - ({}_0^e \Pi(t) {}_0^e \mathbf{X}_1^T - (c + \mu) {}_0^e \mathbf{X}_2^T \\ \quad + \frac{\sigma_1}{\beta} {}_0^e \Pi(t) {}_0^e \mathbf{X}_1^T \mathcal{B}_1(t)) {}_0^e \Theta(t) \simeq 0, \\ \Psi_3^1(t) \triangleq {}_0^e \mathbf{X}_3^T \theta(t) - (c\rho(1-q) {}_0^e \mathbf{X}_2^T + \varphi\bar{\rho} {}_0^e \mathbf{X}_5^T + \alpha_1 {}_0^e \mathbf{X}_4^T \\ \quad - (\gamma_1 + \delta_1 + \mu_1 + \mu) {}_0^e \mathbf{X}_3^T + \sigma_2 {}_0^e \mathbf{X}_4^T \mathcal{B}_2(t)) {}_0^e \Theta(t) \simeq 0, \\ \Psi_4^1(t) \triangleq {}_0^e \mathbf{X}_4^T \theta(t) - (c(1-\rho)(1-q) {}_0^e \mathbf{X}_2^T + \alpha_A {}_0^e \mathbf{X}_5^T \\ \quad - (\alpha_1 + \gamma_A + \mu) {}_0^e \mathbf{X}_4^T - \sigma_2 {}_0^e \mathbf{X}_4^T \mathcal{B}_2(t)) {}_0^e \Theta(t) \simeq 0, \\ \Psi_5^1(t) \triangleq {}_0^e \mathbf{X}_5^T \theta(t) - (cq {}_0^e \mathbf{X}_2^T - (\alpha_A + \varphi\bar{\rho} + \mu) {}_0^e \mathbf{X}_5^T) {}_0^e \Theta(t) \simeq 0, \\ \Psi_6^1(t) \triangleq {}_0^e \mathbf{X}_6^T \theta(t) - (\delta_1 {}_0^e \mathbf{X}_3^T - (\gamma_H + \mu_1 + \mu) {}_0^e \mathbf{X}_6^T \\ \quad - \sigma_3 {}_0^e \mathbf{X}_6^T \mathcal{B}_3(t)) {}_0^e \Theta(t) \simeq 0, \\ \Psi_7^1(t) \triangleq {}_0^e \mathbf{X}_7^T \theta(t) - (\gamma_1 {}_0^e \mathbf{X}_3^T + \gamma_A {}_0^e \mathbf{X}_4^T + \gamma_H {}_0^e \mathbf{X}_6^T \\ \quad - \mu {}_0^e \mathbf{X}_7^T + \sigma_3 {}_0^e \mathbf{X}_6^T \mathcal{B}_3(t)) {}_0^e \Theta(t) \simeq 0, \end{cases} \tag{11}$$

where

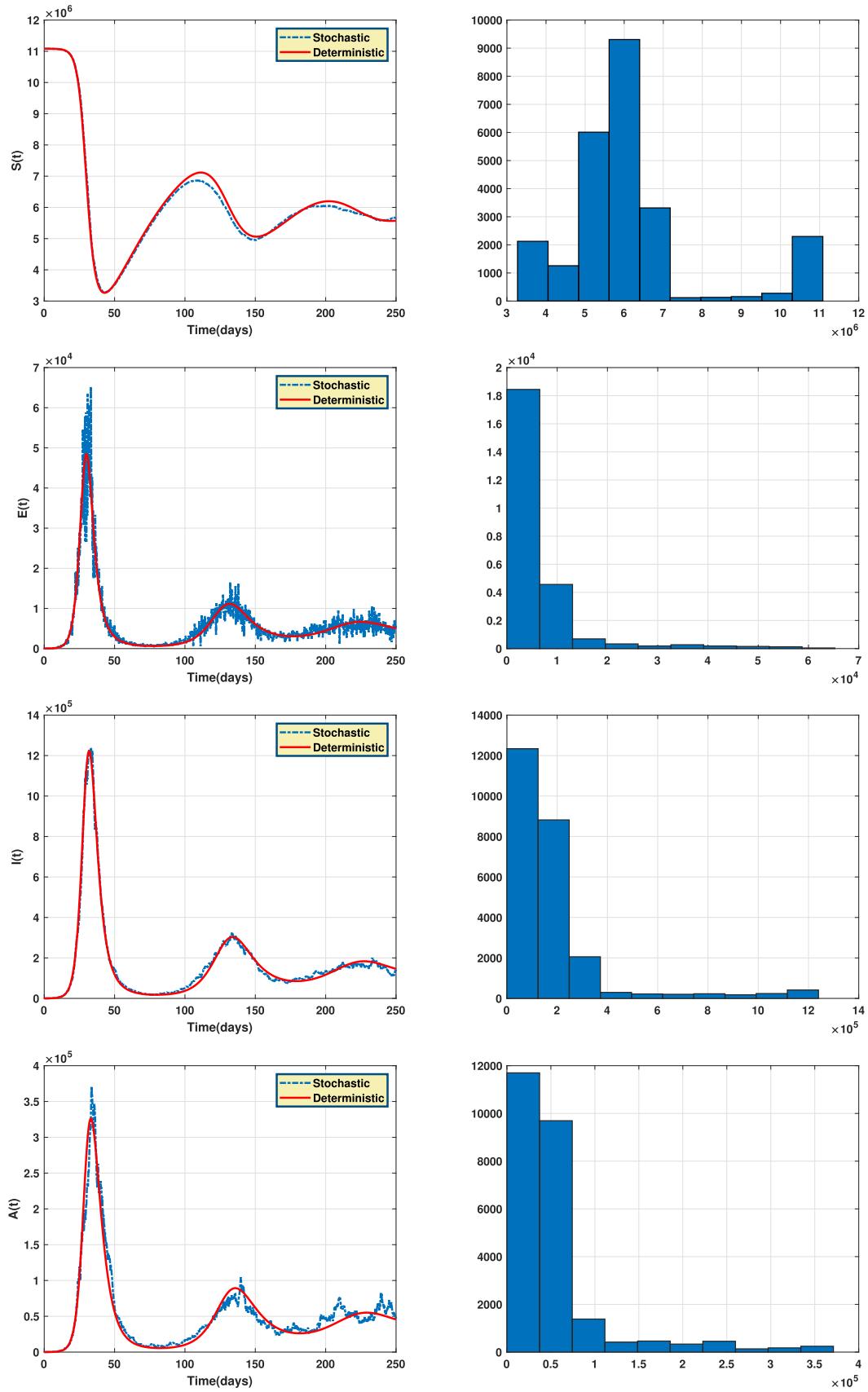
$${}_0^e \tilde{\Theta}(t) := [\partial_t({}_0^e \phi_0(t)), \dots, \partial_t({}_0^e \phi_j(t)), \dots, \partial_t({}_0^e \phi_n(t))]^T,$$

and

$${}_0^e \Pi(t) = \beta(1 - \varepsilon_m c_m) \frac{{}_0^e \mathbf{X}_3^T \theta(t) + \theta {}_0^e \mathbf{X}_4^T \theta(t)}{(\sum_{i=1}^7 {}_0^e \mathbf{X}_i^T \theta(t)) - \theta_4({}_0^e \mathbf{X}_5^T \theta(t) + {}_0^e \mathbf{X}_6^T \theta(t))}.$$

Also, from (10)

$$\begin{aligned} \Phi_1^1 \triangleq & {}_0^e \mathbf{X}_{10}^T \theta(0) - S(0) \simeq 0, & \Phi_2^1 \triangleq & {}_0^e \mathbf{X}_{20}^T \theta(0) - E(0) \simeq 0, \\ \Phi_3^1 \triangleq & {}_0^e \mathbf{X}_{30}^T \theta(0) - I(0) \simeq 0, & \Phi_4^1 \triangleq & {}_0^e \mathbf{X}_{40}^T \theta(0) - A(0) \simeq 0, \\ \Phi_5^1 \triangleq & {}_0^e \mathbf{X}_{50}^T \theta(0) - Q(0) \simeq 0, & \Phi_6^1 \triangleq & {}_0^e \mathbf{X}_{60}^T \theta(0) - H(0) \simeq 0, \\ \Phi_7^1 \triangleq & {}_0^e \mathbf{X}_{70}^T \theta(0) - R(0) \simeq 0. \end{aligned} \tag{12}$$



**Fig. 4.** The trajectories of solution for the stochastic model (6) and the solution of the corresponding deterministic model (left) along with the histograms of frequencies for the populations  $S(t)$ ,  $E(t)$ ,  $I(t)$  and  $A(t)$ (right).



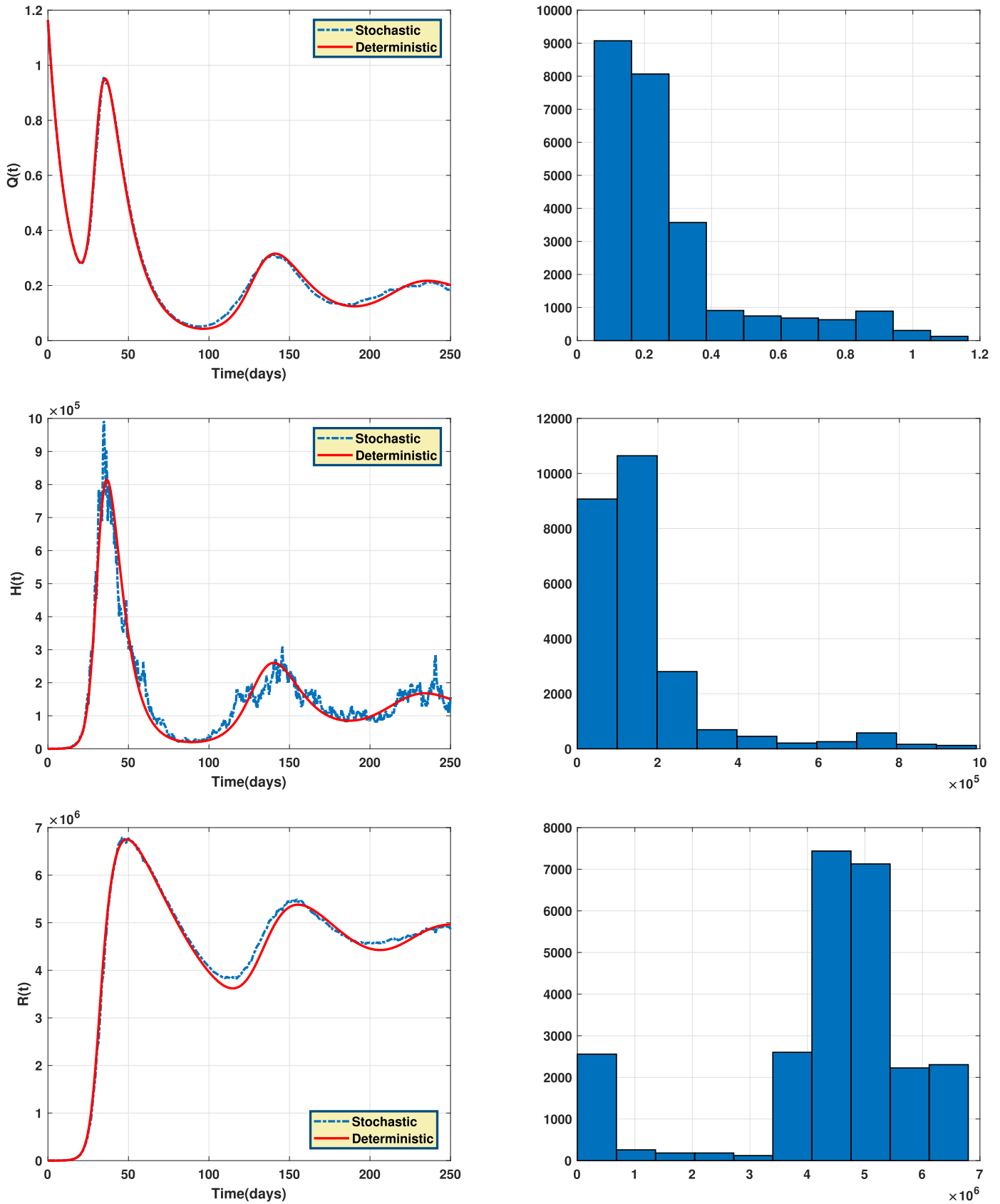


Fig. 5. The trajectories of solution for the stochastic model (6) and the solution of the corresponding deterministic model (left) along with the histograms of frequencies for the populations  $Q(t)$ ,  $H(t)$  and  $R(t)$  (right).

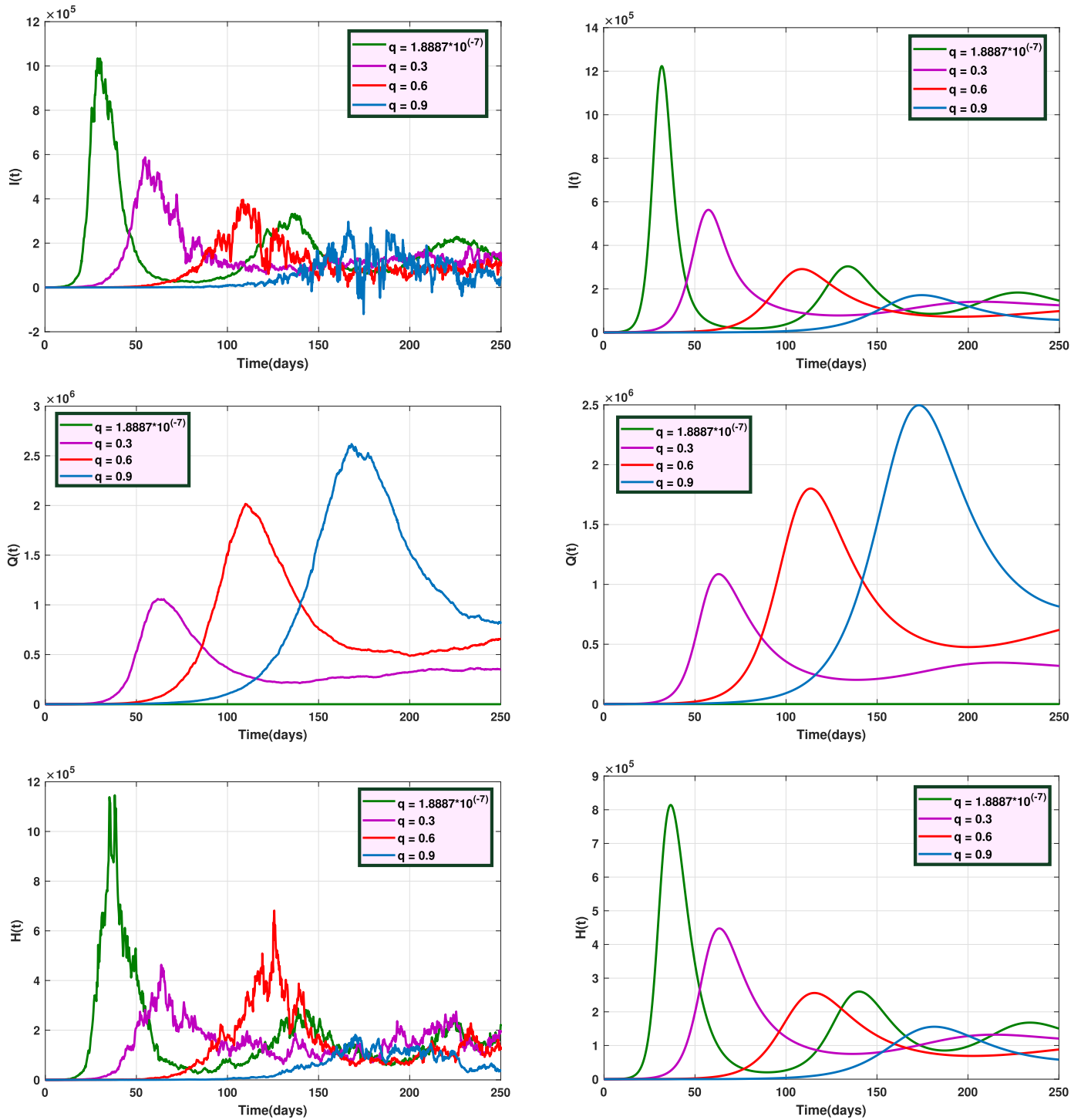


Fig. 6. Graphs of trajectories of  $I(t)$ ,  $Q(t)$  and  $H(t)$  for the stochastic model (left) and the deterministic model (right) with different values of  $q$ .

Let  $t_0^{0,e} = 0$ ,  $t_n^{0,e} = e$ , and  $t_j^{0,e}$ ,  $j = 1, \dots, n-1$ , be the roots of  ${}^0\theta_{n-1}(t)$ . By evaluating Eqs. (11) and (12) at the  $n+1$  collocation points  $t_j^{0,e}$ , a system of  $7(n+1)$  nonlinear algebraic equations is extracted as

$$\begin{cases} \Psi_i^1(t_j^{0,e}) = 0, & i = 1, \dots, 7, \quad j = 1, \dots, n, \\ \Phi_i^1 = 0, & i = 1, \dots, 7. \end{cases} \quad (13)$$

Note that, the part  $\tilde{B}_i(t_j^{0,e})$  in (11) is evaluated by  $B_i(t_j^{0,e}) - B_i(t_{j-1}^{0,e})$ . This system can be solved using a numerical method, such

as Newton's iterative method. Solving this system leads to an approximate solution  ${}_1\mathcal{X}_n^i(t)$ ,  $i = 1, \dots, 7$ , for (6) on the interval  $[0, e]$ .

In general, to find a numerical solution of (6) on the interval  $[(r-1)e, re]$ ,  $r = 2, \dots, \bar{P}$ , suppose  $S(t) \simeq {}_r\mathcal{X}_n^1(t)$ ,  $E(t) \simeq {}_r\mathcal{X}_n^2(t)$ ,  $I(t) \simeq {}_r\mathcal{X}_n^3(t)$ ,  $A(t) \simeq {}_r\mathcal{X}_n^4(t)$ ,  $Q(t) \simeq {}_r\mathcal{X}_n^5(t)$ ,  $H(t) \simeq {}_r\mathcal{X}_n^6(t)$ ,  $R(t) \simeq {}_r\mathcal{X}_n^7(t)$ , in which

$${}_r\mathcal{X}_n^i(t) = \sum_{j=0}^n ({}_{(r-1)e}{}^re\mathbf{x}_j^i) ({}_{(r-1)e}{}^re\theta_j(t)), \quad i = 1, \dots, 7, \quad (14)$$

such that

$$({}_{(r-1)e}{}^re\mathbf{X}_i := [({}_{(r-1)e}{}^re\mathbf{x}_0^i, \dots, ({}_{(r-1)e}{}^re\mathbf{x}_j^i, \dots, ({}_{(r-1)e}{}^re\mathbf{x}_n^i)]^T,$$

**Table 1**  
The parameters values for the considered COVID-19 models.

Parameter	Description	Value	Source
$\Lambda$	Birth rate	$\mu \times N(0)$	[31]
$\beta$	Effective contact rate	0.8648	[22]
$c$	Contact rate	14.781	[31]
$\theta$	Transmission rate	0.5944	[30]
$q$	The quarantine rate	$1.8887 \times 10^{-7}$	[31]
$\delta_1$	Transition rate from E to H	0.13266	[31]
$\gamma$	Recovery rate of I	0.33029	[31]
$\gamma_A$	Recovery rate of A	0.13978	[31]
$\gamma_H$	Recovery rate of H	0.11624	[31]
$\alpha_A$	Movement rate from Q to A	0.059	[22]
$\alpha_1$	Transition rate from A to I	0.078	[22]
$\rho$	Symptoms rate among infected individuals	0.86834	[31]
$c_m$	Proportion of individuals who use masks	0.0546	[22]
$\mu_1$	Mortality rate due to COVID-19	0.01	[30]
$\varepsilon_m$	Efficacy of using masks to prevent catching the infection by S	0.5	[22]
$\theta_q$	Efficacy of quarantine and hospitalization	1	[22]
$\varphi$	Proportion of Q with clinical symptoms	0.05	[29]
$\bar{\rho}$	Movement rate from Q to I with clinical symptoms	0.1259	[29]
$\mu$	Natural death rate	1/69.5	[30]

and

$${}_{(r-1)e}{}^{re}\Theta(t) := [{}_{(r-1)e}{}^{re}\phi_0(t), \dots, {}_{(r-1)e}{}^{re}\phi_j(t), \dots, {}_{(r-1)e}{}^{re}\phi_n(t)]^T.$$

Similarly, by (6) and (14), we get

$$\begin{cases} \Psi_1^r(t) \triangleq {}_{(r-1)e}{}^{re}\mathbf{X}_1^T \tilde{\Theta}(t) - \Lambda + ({}_{(r-1)e}{}^{re}\Pi(t) + \mu)({}_{(r-1)e}{}^{re}\mathbf{X}_1^T \tilde{\Theta}(t)) + \frac{\sigma_1}{\beta} {}_{(r-1)e}{}^{re}\Pi(t) {}_{(r-1)e}{}^{re}\mathbf{X}_1^T \tilde{\Theta}(t) \tilde{B}_1(t) \approx 0, \\ \Psi_2^r(t) \triangleq {}_{(r-1)e}{}^{re}\mathbf{X}_2^T \tilde{\Theta}(t) - ({}_{(r-1)e}{}^{re}\Pi(t) + \mu)({}_{(r-1)e}{}^{re}\mathbf{X}_2^T \tilde{\Theta}(t)) - (c + \mu) {}_{(r-1)e}{}^{re}\mathbf{X}_2^T \tilde{\Theta}(t) + \frac{\sigma_1}{\beta} {}_{(r-1)e}{}^{re}\Pi(t) {}_{(r-1)e}{}^{re}\mathbf{X}_2^T \tilde{\Theta}(t) \tilde{B}_1(t) \approx 0, \\ \Psi_3^r(t) \triangleq {}_{(r-1)e}{}^{re}\mathbf{X}_3^T \tilde{\Theta}(t) - (c\rho(1-q) + \varphi\bar{\rho}) {}_{(r-1)e}{}^{re}\mathbf{X}_3^T \tilde{\Theta}(t) + \alpha_1 {}_{(r-1)e}{}^{re}\mathbf{X}_4^T \tilde{\Theta}(t) - (\gamma_1 + \delta_1 + \mu_1 + \mu) {}_{(r-1)e}{}^{re}\mathbf{X}_3^T \tilde{\Theta}(t) + \sigma_2 {}_{(r-1)e}{}^{re}\mathbf{X}_4^T \tilde{B}_2(t) \approx 0, \\ \Psi_4^r(t) \triangleq {}_{(r-1)e}{}^{re}\mathbf{X}_4^T \tilde{\Theta}(t) - (c(1-\rho)(1-q) + \alpha_A) {}_{(r-1)e}{}^{re}\mathbf{X}_4^T \tilde{\Theta}(t) - (\alpha_1 + \gamma_A + \mu) {}_{(r-1)e}{}^{re}\mathbf{X}_4^T \tilde{\Theta}(t) - \sigma_2 {}_{(r-1)e}{}^{re}\mathbf{X}_4^T \tilde{B}_2(t) \approx 0, \\ \Psi_5^r(t) \triangleq {}_{(r-1)e}{}^{re}\mathbf{X}_5^T \tilde{\Theta}(t) - (cq + \mu) {}_{(r-1)e}{}^{re}\mathbf{X}_5^T \tilde{\Theta}(t) - (\alpha_A + \varphi\bar{\rho} + \mu) {}_{(r-1)e}{}^{re}\mathbf{X}_5^T \tilde{\Theta}(t) \approx 0, \\ \Psi_6^r(t) \triangleq {}_{(r-1)e}{}^{re}\mathbf{X}_6^T \tilde{\Theta}(t) - (\delta_1 + \mu_1 + \mu) {}_{(r-1)e}{}^{re}\mathbf{X}_6^T \tilde{\Theta}(t) - \sigma_3 {}_{(r-1)e}{}^{re}\mathbf{X}_6^T \tilde{B}_3(t) \approx 0, \\ \Psi_7^r(t) \triangleq {}_{(r-1)e}{}^{re}\mathbf{X}_7^T \tilde{\Theta}(t) - (\gamma_H + \mu_1 + \mu) {}_{(r-1)e}{}^{re}\mathbf{X}_7^T \tilde{\Theta}(t) - \gamma_A {}_{(r-1)e}{}^{re}\mathbf{X}_4^T \tilde{\Theta}(t) + \gamma_H {}_{(r-1)e}{}^{re}\mathbf{X}_6^T \tilde{\Theta}(t) - \mu {}_{(r-1)e}{}^{re}\mathbf{X}_7^T \tilde{\Theta}(t) + \sigma_3 {}_{(r-1)e}{}^{re}\mathbf{X}_6^T \tilde{B}_3(t) \approx 0, \end{cases} \quad (15)$$

where

$$\begin{aligned} {}_{(r-1)e}{}^{re}\tilde{\Theta}(t) &:= [\partial_t ({}_{(r-1)e}{}^{re}\phi_0(t)), \dots, \partial_t ({}_{(r-1)e}{}^{re}\phi_j(t)), \dots, \partial_t ({}_{(r-1)e}{}^{re}\phi_n(t))]^T, \\ {}_{(r-1)e}{}^{re}\Pi(t) &= \frac{\beta(1-\varepsilon_m c_m) {}_{(r-1)e}{}^{re}\mathbf{X}_3^T \tilde{\Theta}(t) + \theta {}_{(r-1)e}{}^{re}\mathbf{X}_4^T \tilde{\Theta}(t)}{(\sum_{i=1}^7 {}_{(r-1)e}{}^{re}\mathbf{X}_i^T \tilde{\Theta}(t)) - \theta_q ({}_{(r-1)e}{}^{re}\mathbf{X}_3^T \tilde{\Theta}(t) + {}_{(r-1)e}{}^{re}\mathbf{X}_6^T \tilde{\Theta}(t))} \end{aligned}$$

Let  $t_0^{(x-1)e, re} = (x-1)e$ ,  $t_j^{(x-1)e, re} = re$ , and  $t_j^{(x-1)e, re}$ ,  $j = 1, \dots, n-1$ , be the roots of  ${}_{(x-1)e}{}^{re}\theta_{n-1}(t)$ . By evaluating Eqs. (15) at this collocation points, we have

$$\Psi_i^r(t_j^{(x-1)e, re}) = 0, \quad i = 1, \dots, 7, \quad j = 1, \dots, n. \quad (16)$$

Here,  $\tilde{B}_i(t_j^{(x-1)e, re})$  is estimated by  $B_i(t_j^{(x-1)e, re}) - B_i(t_{j-1}^{(x-1)e, re})$ .

Also, from Eq. (14) at  $t_0^{(x-1)e, re}$ , we get

$$\Phi_i^r \triangleq {}_{(r-1)e}{}^{re}\mathbf{X}_i^T \tilde{\Theta}(t_0^{(x-1)e, re}) - {}_{r-1}\chi_i^i(t_0^{(x-1)e, re}) \approx 0. \quad (17)$$

So, for each step  $r = 2, \dots, \bar{p}$ , the relations (16) and (17) give a system of  $7(n+1)$  nonlinear equations for the unknown coefficients  ${}_{(x-1)e}{}^{re}\mathbf{x}_j^i$ ,  $j = 0, 1, \dots, n$ ,  $i = 1, \dots, 7$ . After solving these systems, the

numerical solutions  $\chi_n^i(t)$ ,  $i = 1, \dots, 7$ , are computed on the interval  $[0, T_{max}]$ , as

$$\chi_n^i(t) = \begin{cases} {}_1\chi_n^i(t), & t \in [0, e], \\ \vdots & \vdots \\ {}_r\chi_n^i(t), & t \in [(r-1)e, re], \\ \vdots & \vdots \\ {}_{\bar{p}}\chi_n^i(t), & t \in [(\bar{p}-1)e, T_{max}], \end{cases} \quad (18)$$

in which  $S(t) \approx \chi_n^1(t)$ ,  $E(t) \approx \chi_n^2(t)$ ,  $I(t) \approx \chi_n^3(t)$ ,  $A(t) \approx \chi_n^4(t)$ ,  $Q(t) \approx \chi_n^5(t)$ ,  $H(t) \approx \chi_n^6(t)$  and  $R(t) \approx \chi_n^7(t)$ .

### 5. Numerical results

In this section, we employ the numerical approach introduced in the previous section to illustrate the obtained results of the proposed models (2) and (6) for COVID-19. These simulations help us to investigate the dynamical behaviors of this pandemic when some safety protocols are observed and random effects are added to the model. To this end, some estimations are considered for the parameters of the model according to some published researches about this disease [22,29-31]. These approximations are displayed in Table 1. Also, we need some initial values for the variables of these models. So, the following initial conditions are supposed:

$$\begin{aligned} S(0) &= 11081739, & E(0) &= 106.2642, & I(0) &= 27.676, \\ A(0) &= 53.839, & Q(0) &= 1.1642, & H(0) &= 1, & R(0) &= 2. \end{aligned} \quad (19)$$

All numerical experiments are performed using Matlab on a PC with a 1.70GHz Core i5-4210 CPU and main memory 8 GB.

Fig. 2 shows the endemic equilibrium  $\mathcal{E}_*$  and the trajectories of the solution for the case  $\mathcal{R}_0 = 0.7730 < 1$  with  $\beta = 0.36$  and the case  $\mathcal{R}_0 = 1.8569 > 1$  with  $\beta = 0.8648$ , when  $\sigma_1 = \sigma_2 = \sigma_3 = 0$ . Fig. 3 displays the reproduction number  $\mathcal{R}_0$  for several values of  $\beta \in [0, 1]$  and  $c_m \in [0, 1]$ , with  $\sigma_1 = \sigma_2 = \sigma_3 = 0$ . Fig. 4 and Fig. 5 display the trajectories of solution for the stochastic model (6) with  $\sigma_1 = 0.15$ ,  $\sigma_2 = 0.32$ ,  $\sigma_3 = 0.26$  and the solution of the corresponding deterministic model along with the histograms of frequencies for  $S(t)$ ,  $E(t)$ ,  $I(t)$ ,  $A(t)$ ,  $Q(t)$ ,  $H(t)$  and  $R(t)$ . These figures confirm that model (6) has a positive solution. Also, from the conditions

$$\frac{\sigma_2^2}{2}(p-1) < \alpha_1 + \gamma_A + \mu,$$

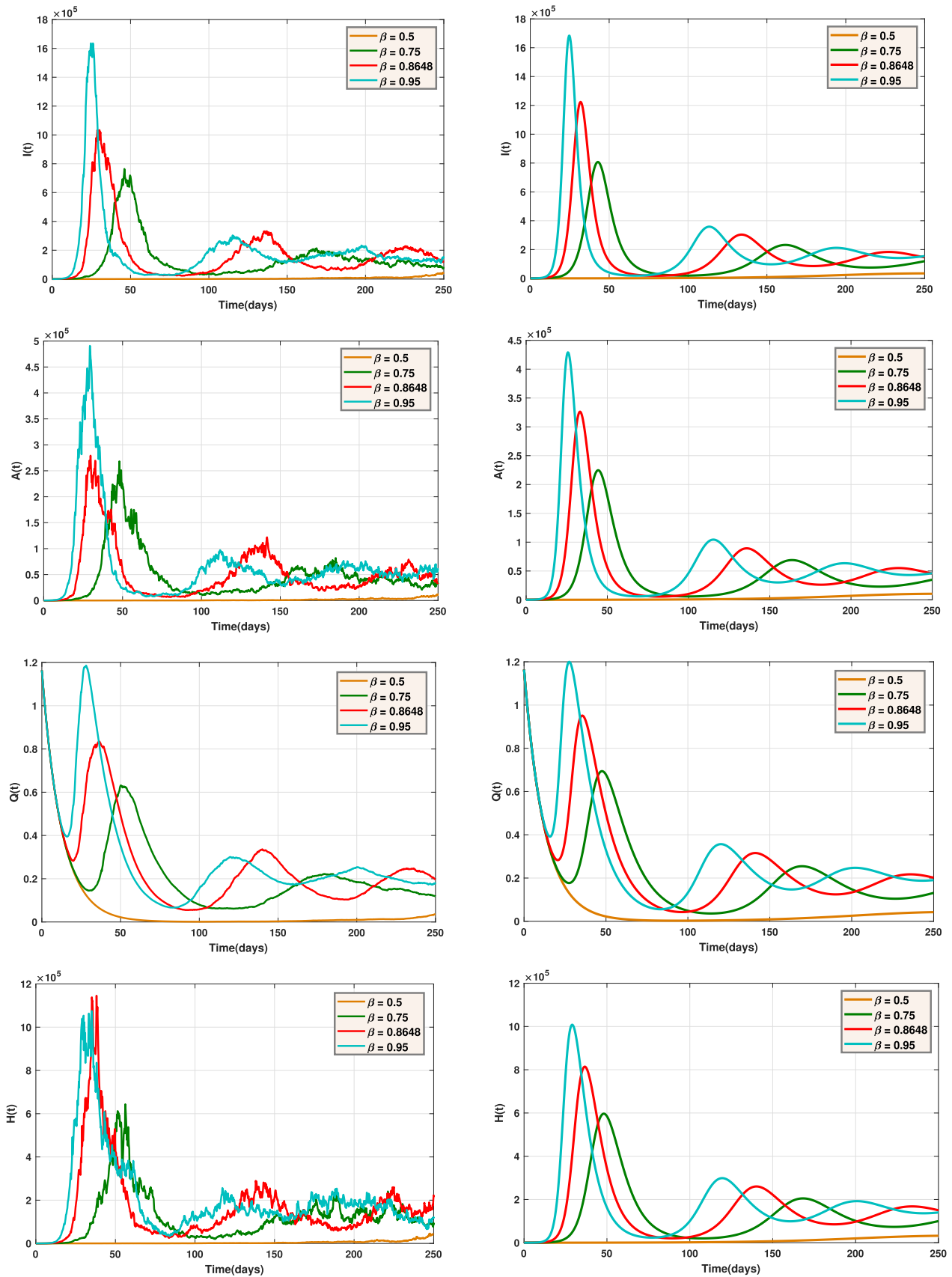


Fig. 7. The trajectories of  $I(t)$ ,  $A(t)$ ,  $Q(t)$  and  $H(t)$  for the stochastic model (left) and the deterministic model (right) with different values of  $\beta$ .

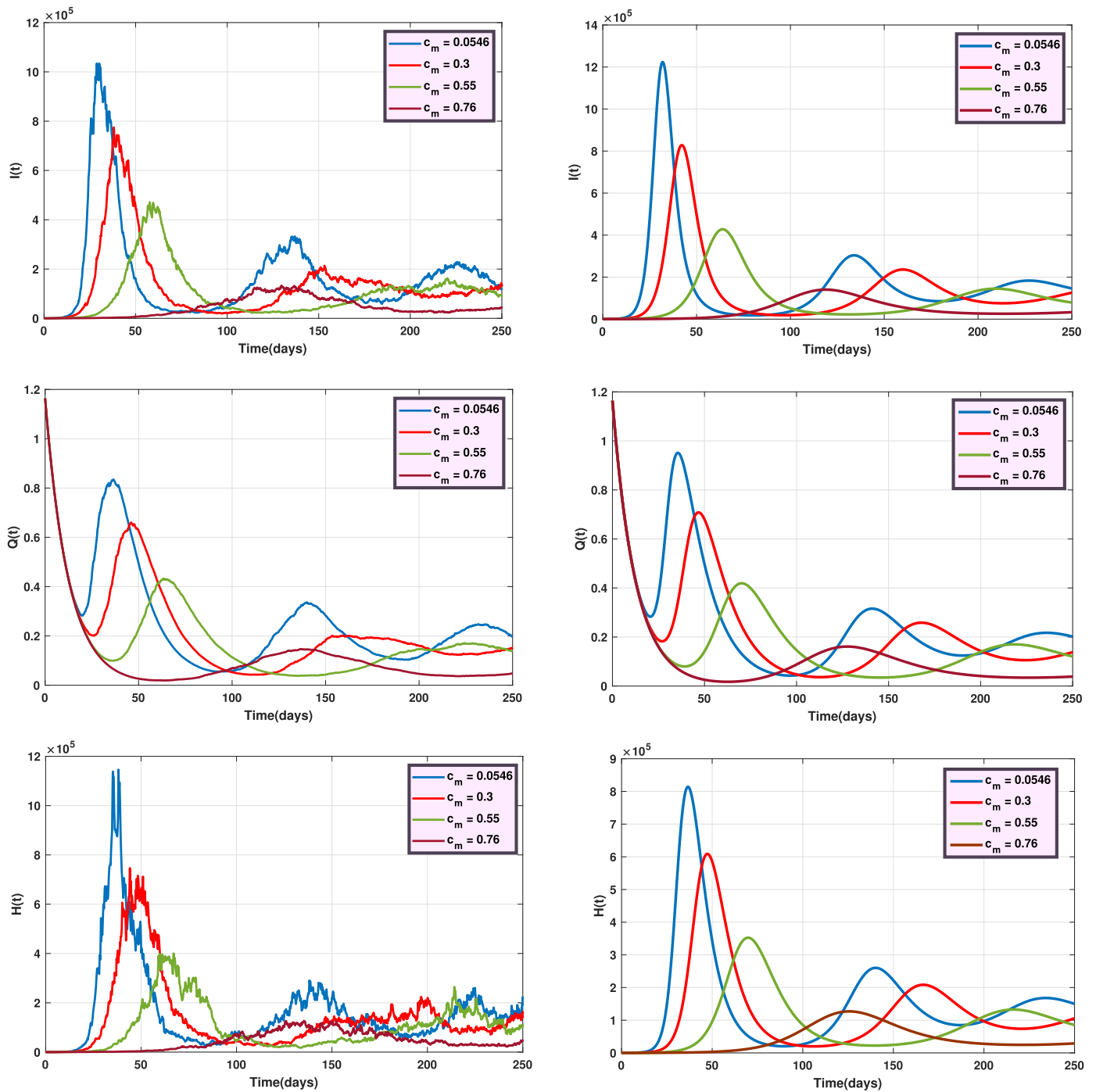


Fig. 8. The trajectories of  $I(t)$ ,  $Q(t)$  and  $H(t)$  for the stochastic model (left) and the deterministic model (right) with different values of  $c_m$ .

and

$$\frac{\sigma_3^2}{2}(p-1) < \gamma_H + \mu_1 + \mu,$$

for  $p \geq 2$  in Theorem 3, we have

$$p = 2 : \quad \frac{\sigma_2^2}{2} = 0.0512 < \alpha_1 + \gamma_A + \mu = 0.2322, \\ \frac{\sigma_3^2}{2} = 0.0338 < \gamma_H + \mu_1 + \mu = 0.1406,$$

$$p = 3 : \quad \sigma_2^2 = 0.1024 < 0.2322, \quad \sigma_3^2 = 0.0676 < 0.1406.$$

Hence, in the case  $p = 2$ ,  $\mathcal{E}^0$  is globally asymptotically stable and in the case  $p = 3$ , the disease-free equilibrium point  $\mathcal{E}^0$  is exponentially 3-stable.

Fig. 6 shows the trajectories of  $I(t)$ ,  $Q(t)$  and  $H(t)$  for the stochastic model when  $\sigma_1 = 0.2$ ,  $\sigma_2 = 0.15$ ,  $\sigma_3 = 0.22$  and the deterministic model, for several values of  $q$ . Fig. 7 displays the trajectories of  $I(t)$ ,  $A(t)$ ,  $Q(t)$  and  $H(t)$  for the stochastic model with  $\sigma_1 = 0.3$ ,  $\sigma_2 = 0.2$ ,  $\sigma_3 = 0.15$  and the deterministic model, with different values of  $\beta$ . Also, Fig. 8 indicates the trajectories of  $I(t)$ ,  $Q(t)$  and  $H(t)$  for the deterministic model and the stochastic model with  $\sigma_1 = 0.15$ ,  $\sigma_2 = 0.24$ ,  $\sigma_3 = 0.36$  for different values of  $c_m$ .

### 6. Conclusion

A stochastic model based on some health protocols such as quarantine, social distancing and wearing masks has outlined to

describe the trend of COVID-19 pandemic in a population. The existence and uniqueness of a positive global solution for the proposed model were proved. Stability analysis of the stochastic model and its special deterministic case have been discussed. In order to simulate the dynamical behaviors of the model, a step-by-step collocation method based on the Legendre polynomials was introduced. Based on the numerical result, it can be seen that considering random effects for the parameters of the model helps to depict a more realistic aspect of the disease development. Also, these results confirm that preventative actions and health strategies have a direct effect on the virus prevalence and control of the pandemic. Considering more public health strategies in the model and studying the impact of underlying diseases or other viral epidemics such as influenza, on the spread of Coronavirus, are some research directions that can be investigated for future works. Also, verifying the frequency of the disease among different age groups may be another useful subject.

### Declaration of Competing Interest

The authors declare no potential conflict of interests.

### CRedit authorship contribution statement

**A. Babaei:** Supervision, Software, Validation, Writing - review & editing. **H. Jafari:** Supervision, Writing - review & editing, Investigation. **S. Banihashemi:** Writing - review & editing, Methodology, Validation, Software. **M. Ahmadi:** Methodology, Validation, Software.

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