



**Applicable Analysis** An International Journal

ISSN: 0003-6811 (Print) 1563-504X (Online) Journal homepage: http://www.tandfonline.com/loi/gapa20

# Stability of a stochastic model for HIV-1 dynamics within a host

Leonid Shaikhet & Andrei Korobeinikov

To cite this article: Leonid Shaikhet & Andrei Korobeinikov (2016) Stability of a stochastic model for HIV-1 dynamics within a host, Applicable Analysis, 95:6, 1228-1238, DOI: 10.1080/00036811.2015.1058363

To link to this article: http://dx.doi.org/10.1080/00036811.2015.1058363

	•	(	<u>}</u>
1			
1			
-1			

Published online: 22 Jun 2015.



Submit your article to this journal

Article views: 103



View related articles



View Crossmark data 🗹



Full Terms & Conditions of access and use can be found at http://www.tandfonline.com/action/journalInformation?journalCode=gapa20



# Stability of a stochastic model for HIV-1 dynamics within a host

Leonid Shaikhet<sup>a</sup> and Andrei Korobeinikov<sup>bc\*</sup>

<sup>a</sup> Department of Higher Mathematics, Donetsk State University of Management, Chelyuskintsev Street 163a, Donetsk 83015, Ukraine; <sup>b</sup>Centre de Recerca Matemática, Campus de Bellaterra, Edifici C, 08193 Bellaterra, Barcelona, Spain; <sup>c</sup>Departament de Matemátiques, Universitat Autónoma de Barcelona, 08193 Bellaterra, Barcelona, Spain

Communicated by M. Yamamoto

(Received 16 May 2014; accepted 1 June 2015)

We consider a stochastically perturbed Nowak–May model of virus dynamics within a host. Using the direct Lyapunov method, we found sufficient conditions for the stability in probability of equilibrium states of this model.

**Keywords:** virus dynamics; stochastic model; direct Lyapunov method; Lyapunov function; equilibrium state

AMS Subject Classifications: Primary: 92D30; Secondary: 34D20; 60H10

## 1. Background

During the last decade, a significant advance was made in the global analysis of mathematical models in epidemiology and viral dynamics. This advance is based on an application of the direct Lyapunov method and follows the initial breakthrough in [1,2]. In these paper, the so-called Volterra–Lyapunov function of the form

$$V(x_1,...,x_n) = \sum_{i=1}^n A_i \left( x_i(t) - x_i^* \ln x_i(t) \right),$$

where  $A_i$  are positive constants which have to be defined,  $x_i(t)$  are the phase variables, and  $x_i^*$  are coordinates of the equilibrium state to be studied, was first time successfully applied to a three-dimensional SEIR model. (It is worthy of mentioning that application of this function to two-dimensional models had been well developed by this time.) This result initiated a vast literature, which currently includes a few thousands of items. This technique was applied by many other authors to a variety of models in mathematical biology and epidemiology, and subsequently extended to the models with an arbitrary number of subpopulations and compartments,[3–7] models with nonlinear functional responses,[8–11] models with time delays [12,13] and models with distributed subpopulations.[14–17]

However, all these results have a certain drawback, as they are dealing with comparatively simple deterministic models which ability to reflect the complexity of real-life

<sup>\*</sup>Corresponding author. Email: akorobeinikov@crm.cat

<sup>© 2015</sup> Taylor & Francis

biological systems is rather limited. Processes in biology are extremely complex and involve a large number of factors; impacts of these factors are often not known in details. Moreover, details of the interactions between the major agents are usually also known with a large degree of uncertainty. In the deterministic models framework, this complexity can be to some extend captured by further increase of the model order, by introduction of delays and by distributed parameters. However, the majority of factors, which are involved in real-life processes or have influence on these, are not known in sufficient details. In such a situation, the use of stochastic models, where these complexity and uncertainty can be captured by stochastic perturbations, appears to be a natural choice. Moreover, analysing the impact which stochastic perturbation can have the dynamics of deterministic models, can justify the results obtained for these deterministic models. However, the analysis of stochastic models is a considerably more challenging task, and it is hardly surprising, therefore, that just a few dozens of publications, dealing with the analysis of stochastic models in epidemiology and viral dynamics, appeared for the last decade.

In this notice, we consider a stochastically perturbed model of virus dynamics within a host; the deterministic version of this model was suggested by Nowak and May [17] in order to describe the dynamics of HIV-1 in vivo. Our objective is to establish sufficient conditions of the stability in probability [18] for equilibrium states of this model. To address this problem, we apply the direct Lyapunov method.

The Nowak–May model of virus dynamics comprises three variable quantities, namely concentrations of the susceptible target cells, x(t), the infected cells, y(t) and free virus particles, v(t), respectively. The model postulates that there is a constant influx of the susceptible target cells with a rate  $\lambda$ , that the susceptible cells are infected by free virus particles with a bilinear incidence rate  $\beta x(t)v(t)$ , and that the infected cells produce free virus particles with a rate ky(t); average life spans of the susceptible cells (in absence of the virus), the infected cells and free virus particles are 1/m, 1/a and 1/u, respectively. Under these assumptions, the model is represented by the following system of ordinary differential equations:

$$\dot{x}(t) = \lambda - mx(t) - \beta x(t)v(t),$$
  

$$\dot{y}(t) = \beta x(t)v(t) - ay(t),$$
  

$$\dot{v}(t) = ky(t) - uv(t).$$
(1.1)

Equilibrium states of this model satisfy the following system of algebraic equations:

$$\lambda = mx + \beta xv,$$
  

$$ay = \beta xv,$$
  

$$ky = uv.$$
(1.2)

It is easy to see that the model can have at most two equilibrium states, namely an infection-free equilibrium state  $E_0 = (x_0, y_0, z_0)$ , where

$$x_0 = \frac{\lambda}{m}, \qquad y_0 = 0, \qquad v_0 = 0,$$
 (1.3)

and a positive equilibrium state  $E^* = (x^*, y^*, v^*)$ , where

$$x^* = \frac{au}{k\beta}, \qquad y^* = \frac{\lambda}{a} - \frac{mu}{k\beta}, \qquad v^* = \frac{\lambda k}{au} - \frac{m}{\beta}.$$
 (1.4)

The positive equilibrium state  $E^*$  exists if the condition

$$R_0 = \frac{\lambda k\beta}{amu} > 1, \tag{1.5}$$

holds. Here,  $R_0$  is called the basic reproduction number of virus.

The properties of the Nowak–May model are well studied. Specifically, it was proved that if  $R_0 \le 1$  holds, then the infection-free equilibrium  $E_0$  (which is the only equilibrium state of the model in this case) is globally asymptotically stable, whereas if  $R_0 > 1$  holds, then  $E_0$  is a saddle point, and the positive equilibrium state  $E^*$  exists and is globally asymptotically stable (in  $\mathbb{R}^3_+$ , as *x*-axis is an invariant set of the model and the stable manifold of  $E_0$ ).[2,19]

## 2. A stochastically perturbed model

Let us now assume that system (1.1) is stochastically perturbed by white noises, and that magnitudes of perturbations are proportional to the deviation of a current state (x(t), y(t), v(t)) from an equilibrium point. Thus, for the positive equilibrium state  $E^*$ , the stochastically perturbed system takes the form

$$\dot{x}(t) = \lambda - mx(t) - \beta x(t)v(t) + \sigma_1(x(t) - x^*)\dot{w}_1(t),$$
  

$$\dot{y}(t) = \beta x(t)v(t) - ay(t) + \sigma_2(y(t) - y^*)\dot{w}_2(t),$$
  

$$\dot{v}(t) = ky(t) - uv(t) + \sigma_3(v(t) - v^*)\dot{w}_3(t),$$
(2.1)

where  $\sigma_1$ ,  $\sigma_2$  and  $\sigma_3$  are positive constants, and  $w_1(t)$ ,  $w_2(t)$  and  $w_3(t)$  are mutually independent Wiener processes. An advantage of this approach is that for the stochastic perturbations of such a type the equilibrium point  $(x^*, y^*, v^*)$  of system (2.1) coincides with the equilibrium point  $E^*$  of system (1.1).

Stochastic perturbations of this form were for the first time proposed in [20] for SIR epidemic model and later this idea was successfully applied by different authors to different mathematical models, described by differential equations (see [18]) and references therein) by finite difference equations with discrete and continuous time,[21] by partial differential equations.[22] The technique which we use further in this paper for the analysis of system (2.1) is similar to that in [18,20–22].

A substitution

$$x(t) = x_1(t) + x^*,$$
  $y(t) = x_2(t) + y^*,$   $v(t) = x_3(t) + v^*$ 

transforms system (2.1) to the form

$$\dot{x}_1(t) = \lambda - m(x_1(t) + x^*) - \beta(x_1(t) + x^*)(x_3(t) + v^*) + \sigma_1 x_1(t) \dot{w}_1(t),$$
  
$$\dot{x}_2(t) = \beta(x_1(t) + x^*)(x_3(t) + v^*) - a(x_2(t) + y^*) + \sigma_2 x_2(t) \dot{w}_2(t),$$
  
$$\dot{x}_3(t) = k(x_2(t) + y^*) - u(x_3(t) + v^*) + \sigma_3 x_3(t) \dot{w}_3(t),$$

or, using (1.2), to the form

$$\dot{x}_1(t) = -(m + \beta v^*)x_1(t) - \beta x^* x_3(t) - \beta x_1(t)x_3(t) + \sigma_1 x_1(t)\dot{w}_1(t),$$
  

$$\dot{x}_2(t) = \beta (v^* x_1(t) + x^* x_3(t) + x_1(t)x_3(t)) - ax_2(t) + \sigma_2 x_2(t)\dot{w}_2(t),$$
  

$$\dot{x}_3(t) = kx_2(t) - ux_3(t) + \sigma_3 x_3(t)\dot{w}_3(t).$$
(2.2)

1230

The origin of the system (2.2) phase space corresponds to the positive equilibrium state  $E^*$  of system (2.1) and is an equilibrium state of this system. Our objective is to establish the stability of the origin.

Further, we will also consider the linear part of system (2.2) in the form

$$\dot{y}_1(t) = -(m + \beta v^*)y_1(t) - \beta x^* y_3(t) + \sigma_1 y_1(t)\dot{w}_1(t),$$
  

$$\dot{y}_2(t) = \beta (v^* y_1(t) + x^* y_3(t)) - a y_2(t) + \sigma_2 y_2(t)\dot{w}_2(t),$$
  

$$\dot{y}_3(t) = k y_2(t) - u y_3(t) + \sigma_3 y_3(t)\dot{w}_3(t).$$
(2.3)

Note that the order of nonlinearity of the system (2.2) is higher than one, and hence sufficient conditions for the asymptotic mean square stability (Definition A.1) of the trivial solution of the system (2.3) are, at the same time, the sufficient conditions for the stability in probability (Definition A.1) for the trivial solution of the system (2.2) (Remark A.1).

## 3. Stability for the positive equilibrium state *E*

In order to establish conditions for the stability of positive equilibrium state  $E^*$ , we represent the linear system (2.3) in the matrix form

$$\dot{y}(t) = Ay(t) + B(y(t))\dot{w}(t);$$
(3.1)

here

$$y(t) = \begin{pmatrix} y_1(t) \\ y_2(t) \\ y_3(t) \end{pmatrix}, \quad A = \begin{pmatrix} -(m + \beta v^*) & 0 & -\beta x^* \\ \beta v^* & -a & \beta x^* \\ 0 & k & -u \end{pmatrix},$$
$$B(y(t)) = \begin{pmatrix} \sigma_1 y_1(t) & 0 & 0 \\ 0 & \sigma_2 y_2(t) & 0 \\ 0 & 0 & \sigma_3 y_3(t) \end{pmatrix}, \quad w(t) = \begin{pmatrix} w_1(t) \\ w_2(t) \\ w_3(t) \end{pmatrix}.$$

It is easy to see that, if  $v^* > 0$  (that is, if  $R_0 > 1$ ) holds, then matrix A satisfies the Routh– Hurwitz criterion for stability (see [23], p.197). Indeed, for coefficients of the characteristic equation

$$\lambda^3 - S_1\lambda^2 + S_2\lambda - S_3 = 0,$$

the inequalities

$$S_{1} = trA = -(m + \beta v^{*} + a + u) < 0,$$
  

$$S_{2} = (m + \beta v^{*})(a + u) > 0,$$
  

$$S_{3} = \det A = -au\beta v^{*} < 0,$$
  

$$S_{3} > S_{1}S_{2}$$

hold (we used equalities (1.2) here). Therefore, for a positive definite matrix

$$Q = \begin{pmatrix} q_1 & 0 & 0\\ 0 & q_2 & 0\\ 0 & 0 & 1 \end{pmatrix}, \qquad q_1 > 0, \qquad q_2 > 0,$$

there exists a positive definite solution  $P = (p_{ij})$  of the Lyapunov matrix equation

$$A'P + PA = -Q. \tag{3.2}$$

*Remark 3.1* The diagonal elements  $p_{ii}$ , i = 1, 2, 3, of the solution *P* to the matrix Equation (3.2) are defined [18] as

$$p_{ii} = \frac{1}{2\Delta_3} \sum_{r=0}^{2} \gamma_i^{(r)} \Delta_{1,r+1}.$$
(3.3)

Here,

$$\Delta_3 = \begin{vmatrix} -S_1 & -S_3 & 0\\ 1 & S_2 & 0\\ 0 & -S_1 & -S_3 \end{vmatrix} = (S_1 S_2 - S_3) S_3 > 0$$

is the determinant of Hurwitz matrix;  $\Delta_{1,r+1}$  is the algebraic adjunct of the element of the first line and (r + 1)th column of the determinant  $\Delta_3$ , that is,  $\Delta_{11} = -S_2S_3$ ,  $\Delta_{12} = S_3$  and  $\Delta_{13} = -S_1$  and  $\gamma_i^{(r)}$  are defined by the identity

$$\sum_{k=1}^{3} q_k D_{ik}(\lambda) D_{ik}(-\lambda) \equiv \sum_{r=0}^{2} \gamma_i^{(r)} \lambda^{2(2-r)},$$
(3.4)

where  $q_i$  are the elements of the matrix Q, that is,  $q_1 > 0$ ,  $q_2 > 0$  and  $q_3 = 1$ , and  $D_{ik}(\lambda)$  are the algebraic adjuncts of the determinant

$$D(\lambda) = \begin{vmatrix} a_{11} - \lambda & a_{12} & a_{13} \\ a_{21} & a_{22} - \lambda & a_{23} \\ a_{31} & a_{32} & a_{33} - \lambda \end{vmatrix}.$$

From (3.3), (3.4) it follows that the diagonal elements of the solution P of the matrix Equation (3.2) can be represented in the form

$$p_{ii} = p_{ii}^{(1)} q_1 + p_{ii}^{(2)} q_2 + p_{ii}^{(3)}, \qquad i = 1, 2, 3.$$
(3.5)

Sufficient stability conditions for the trivial solution of system (2.3) are given by following Theorem:

THEOREM 3.1 Let  $\sigma_i$ , i = 1, 2, 3, satisfy the conditions

$$\sigma_{1}^{2} < \left(p_{11}^{(1)}\right)^{-1}, \qquad \sigma_{2}^{2} < \left(p_{22}^{(2)} + \frac{\sigma_{1}^{2} p_{11}^{(2)} p_{22}^{(1)}}{1 - \sigma_{1}^{2} p_{11}^{(1)}}\right)^{-1}, \\ \sigma_{3}^{2} < \left(p_{33}^{(3)} + \frac{\sigma_{1}^{2} p_{33}^{(1)} \left[\sigma_{2}^{2} p_{11}^{(2)} p_{22}^{(3)} + p_{11}^{(3)} \left(1 - \sigma_{2}^{2} p_{22}^{(2)}\right)\right]}{\left(1 - \sigma_{1}^{2} p_{11}^{(1)}\right) \left(1 - \sigma_{2}^{2} p_{22}^{(2)}\right) - \sigma_{1}^{2} \sigma_{2}^{2} p_{11}^{(2)} p_{22}^{(1)}} \\ + \frac{\sigma_{2}^{2} p_{33}^{(2)} \left[\sigma_{1}^{2} p_{11}^{(3)} p_{22}^{(1)} + p_{22}^{(3)} \left(1 - \sigma_{1}^{2} p_{11}^{(1)}\right)\right]}{\left(1 - \sigma_{1}^{2} p_{11}^{(1)}\right) \left(1 - \sigma_{2}^{2} p_{22}^{(2)}\right) - \sigma_{1}^{2} \sigma_{2}^{2} p_{11}^{(2)} p_{22}^{(1)}}\right)^{-1}.$$
(3.6)

Then, the trivial solution of system (2.3) is asymptotically mean square stable.

*Proof* Let *L* be the generator (see Appendix 1) of the Equation (3.1),  $P = (p_{ij})$  be the solution of the matrix Equation (3.2), and V = y' P y. Then,

$$LV = y'(t)(A'P + PA)y(t) + \sum_{i=1}^{3} \sigma_i^2 p_{ii} y_i^2(t)$$
  
=  $\left(-q_1 + \sigma_1^2 p_{11}\right) y_1^2(t) + \left(-q_2 + \sigma_2^2 p_{22}\right) y_2^2(t) + \left(-1 + \sigma_3^2 p_{33}\right) y_3^2(t),$ 

and hence the sufficient conditions for the asymptotic mean square stability of the trivial solution of Equation (3.1) are (see Theorem A.1)

 $\sigma_1^2 p_{11} < q_1, \qquad \sigma_2^2 p_{22} < q_2, \qquad \sigma_3^2 p_{33} < 1.$  (3.7)

Using (3.5), we can rewrite conditions (3.7) in the form

$$\sigma_1^2 \left( p_{11}^{(1)} q_1 + p_{11}^{(2)} q_2 + p_{11}^{(3)} \right) < q_1, \tag{3.8}$$

$$\sigma_2^2 \left( p_{22}^{(1)} q_1 + p_{22}^{(2)} q_2 + p_{22}^{(3)} \right) < q_2, \tag{3.9}$$

$$\sigma_3^2 \left( p_{33}^{(1)} q_1 + p_{33}^{(2)} q_2 + p_{33}^{(3)} \right) < 1.$$
(3.10)

It is easy to see that Theorem hypotheses (3.6) imply that (3.8)–(3.10) hold for some positive  $q_1$  and  $q_2$ . Indeed, combining inequalities (3.8) and (3.9), we get

$$0 < \frac{\sigma_2^2 \left( p_{22}^{(1)} q_1 + p_{22}^{(3)} \right)}{1 - \sigma_2^2 p_{22}^{(2)}} < q_2 < \frac{\left( 1 - \sigma_1^2 p_{11}^{(1)} \right) q_1 - \sigma_1^2 p_{11}^{(3)}}{\sigma_1^2 p_{11}^{(2)}}.$$
 (3.11)

Hence, if there exists  $q_1 > 0$  such that the inequality

$$0 < \frac{\sigma_2^2 \left( p_{22}^{(1)} q_1 + p_{22}^{(3)} \right)}{1 - \sigma_2^2 p_{22}^{(2)}} < \frac{\left( 1 - \sigma_1^2 p_{11}^{(1)} \right) q_1 - \sigma_1^2 p_{11}^{(3)}}{\sigma_1^2 p_{11}^{(2)}}$$
(3.12)

holds, then there also exists  $q_2 > 0$  such that (3.11) holds. Furthermore, from (3.12) it follows that

$$q_{1} > \frac{\sigma_{1}^{2} \left[ \sigma_{2}^{2} p_{11}^{(2)} p_{22}^{(3)} + p_{11}^{(3)} \left( 1 - \sigma_{2}^{2} p_{22}^{(2)} \right) \right]}{\left( 1 - \sigma_{1}^{2} p_{11}^{(1)} \right) \left( 1 - \sigma_{2}^{2} p_{22}^{(2)} \right) - \sigma_{1}^{2} \sigma_{2}^{2} p_{11}^{(2)} p_{22}^{(1)}}.$$
(3.13)

Likewise, inequalities (3.8) and (3.9) yield

$$0 < \frac{\sigma_1^2 \left( p_{11}^{(2)} q_2 + p_{11}^{(3)} \right)}{1 - \sigma_1^2 p_{11}^{(1)}} < q_1 < \frac{\left( 1 - \sigma_2^2 p_{22}^{(2)} \right) q_2 - \sigma_2^2 p_{22}^{(3)}}{\sigma_2^2 p_{22}^{(1)}}, \tag{3.14}$$

and hence the existence of  $q_2 > 0$ , such that the inequality

$$0 < \frac{\sigma_1^2 \left( p_{11}^{(2)} q_2 + p_{11}^{(3)} \right)}{1 - \sigma_1^2 p_{11}^{(1)}} < \frac{\left( 1 - \sigma_2^2 p_{22}^{(2)} \right) q_2 - \sigma_2^2 p_{22}^{(3)}}{\sigma_2^2 p_{22}^{(1)}}$$
(3.15)

holds, also ensures the existence of  $q_1 > 0$  such that (3.14) holds. Moreover, from (3.15), it follows that

$$q_{2} > \frac{\sigma_{2}^{2} \left[ \sigma_{1}^{2} p_{11}^{(3)} p_{22}^{(1)} + p_{22}^{(3)} \left( 1 - \sigma_{1}^{2} p_{11}^{(1)} \right) \right]}{\left( 1 - \sigma_{1}^{2} p_{11}^{(1)} \right) \left( 1 - \sigma_{2}^{2} p_{22}^{(2)} \right) - \sigma_{1}^{2} \sigma_{2}^{2} p_{11}^{(2)} p_{22}^{(1)}}.$$
(3.16)

Theorem hypothesis, and specifically the first and the second inequalities in (3.6), guaranties that the right-hand parts in (3.13) and (3.16) are positive. Furthermore, by (3.13), (3.16) and the third condition in (3.6), inequality (3.10) holds as well. That is, Theorem hypothesis (3.6) ensure the existence of  $q_1 > 0$  and  $q_2 > 0$  such that conditions (3.8)–(3.10) hold, and, thereby, the asymptotic mean square stability of the trivial solution of linear Equation (2.3). The proof is now completed.

COROLLARY 3.1 Under Theorem 3.1.1 hypothesis, the trivial solution of nonlinear system (2.2), or, what is the same, of the positive equilibrium state  $E^*$  of system (2.1), is stable in probability.

The proof follows from Theorems A.1, A.2 and Remark A.1 (see Appendix 1), since the order of nonlinearity of the system (2.2) is higher than one.

*Remark 3.2* Note that if all conditions (3.7) do not hold then  $LV \ge 0$  and the trivial solution of the system (2.3) cannot be asymptotically mean square stable.[18]

#### 4. Stability of the infection-free equilibrium $E_0$

For the infection-free equilibrium state  $E_0$ , systems (2.1)–(2.3), respectively, have the forms

$$\dot{x}(t) = \lambda - mx(t) - \beta x(t)v(t) + \sigma_1(x(t) - x_0)\dot{w}_1(t), \dot{y}(t) = \beta x(t)v(t) - ay(t) + \sigma_2 y(t)\dot{w}_2(t), \dot{v}(t) = ky(t) - uv(t) + \sigma_3 v(t)\dot{w}_3(t);$$
(4.1)

$$\dot{x}_1(t) = -mx_1(t) - \beta x_0 x_3(t) - \beta x_1(t) x_3(t) + \sigma_1 x_1(t) \dot{w}_1(t),$$
  

$$\dot{x}_2(t) = \beta (x_0 x_3(t) + x_1(t) x_3(t)) - a x_2(t) + \sigma_2 x_2(t) \dot{w}_2(t),$$
  

$$\dot{x}_3(t) = k x_2(t) - u x_3(t) + \sigma_3 x_3(t) \dot{w}_3(t),$$
(4.2)

and

$$\dot{y}_{1}(t) = -my_{1}(t) - \beta x_{0}y_{3}(t) + \sigma_{1}y_{1}(t)\dot{w}_{1}(t),$$
  

$$\dot{y}_{2}(t) = -ay_{2}(t) + \beta x_{0}y_{3}(t) + \sigma_{2}y_{2}(t)\dot{w}_{2}(t),$$
  

$$\dot{y}_{3}(t) = ky_{2}(t) - uy_{3}(t) + \sigma_{3}y_{3}(t)\dot{w}_{3}(t).$$
(4.3)

For these systems, conditions for the asymptotic mean square stability of the trivial solution of system (4.3) at the same time are the conditions for the stability in probability of the trivial solution of system (4.2) and for the stability in probability of the infection-free equilibrium state  $E_0$  of system (4.1).

Conditions of the stability are given by following Theorem:

THEOREM 4.1 Let  $R_0 < 1$  holds, and  $\sigma_i$ , i = 1, 2, 3, satisfy the conditions

$$\delta_1 < m, \qquad \delta_2 < \frac{|tr(A)|\det(A)}{A_2}, \qquad \delta_3 < \frac{|tr(A)|\det(A) - A_2\delta_2}{A_1 - |tr(A)|\delta_2}, \qquad (4.4)$$

where

$$\delta_{i} = \frac{1}{2}\sigma_{i}^{2}, \quad i = 1, 2, 3,$$

$$A = \begin{pmatrix} -a \ \beta x_{0} \\ k \ -u \end{pmatrix}, \quad tr(A) = -(a+u) < 0,$$

$$\det(A) = au - k\beta x_{0} = au - k\beta\lambda m^{-1} = au(1-R_{0}) > 0,$$

$$A_{1} = \det(A) + a^{2}, \quad A_{2} = \det(A) + u^{2}.$$

Then, the trivial solution of the system (4.3) is asymptotically mean square stable.

**Proof** We note that the second and the third equations of (4.3) are independent of  $y_1(t)$ , and hence the system of two equation  $(y_2(t), y_3(t))$  can be considered separately. Sufficient conditions for the asymptotic mean square stability of the trivial solutions for systems of such a type are well known (for instance, see [18]). Theorem hypothesis (4.4) ensures that these conditions are held. The conditions (4.4) also imply the stability in probability of the equilibrium point (1.3) of the system (4.1).

#### 5. Conclusion

The concept of stochastic modelling allows to capture, to some extent, the complexity of biological processes and to imitate the impacts of unavoidable in the real-life uncertainties. This consideration motivates the growing interest to stochastic modelling in mathematical biology. In this paper, we considered a stochastic version of the Nowak–May model of virus dynamics within a host, which was originally suggested to describe HIV-1 dynamics. Using the direct Lyapunov method, we found sufficient conditions for the stability in probability for the equilibrium states of this model.

It is noteworthy that the approach used in this paper can be applied to a wider variety of the stochastic models.

#### **Disclosure statement**

No potential conflict of interest was reported by the authors.

#### Funding

A. Korobeinikov is supported by the Ministry of Science and Innovation of Spain via Ramón y Cajal Fellowship [RYC-2011-08061] and [grant number MTM2011-29342], by CONACYT (Mexico) via [grant number N219614], and by AGAUR (Generalitat de Catalunya) via [grant number 2014SGR-1307].

# References

- Korobeinikov A. Lyapunov functions and global properties for SEIR and SEIS epidemic models. Math. Med. Biol. 2004;21:75–83.
- [2] Korobeinikov A. Global properties of basic virus dynamics models. Bull. Math. Biol. 2004;66:879–883.
- [3] Adda P, Dimi JL, Iggidr A, Kamgang JC, Sallet G, Tewa JJ. General models of host-parasite systems. Global analysis. Disc. Cont. Dyn. Syst. B. 2007;8:1–17.
- [4] Fall A, Iggidr A, Sallet G, Tewa JJ. Epidemiological models and Lyapunov functions. Math. Model. Nat. Phenom. 2007;2:62–83.
- [5] Korobeinikov A. Global properties of SIR and SEIR epidemic models with multiple parallel infectious stages. Bull. Math. Biol. 2009;71:75–83.
- [6] Melnik AV, Korobeinikov A. Global asymptotic properties of staged models with multiple progression pathways for infectious diseases. Math. Biosci. Eng. 2011;8:1019–1034.
- [7] Okuonghae D, Korobeinikov A. Dynamics of tuberculosis: the effect of direct observation therapy strategy (DOTS) in Nigeria. Math. Model. Nat. Phenom. 2006;2:99–111.
- [8] Korobeinikov A. Lyapunov functions and global stability for SIR and SIRS epidemiological models with non-linear transmission. Bull. Math. Biol. 2006;68:615–626.
- Korobeinikov A. Global properties of infectious disease models with non-linear incidence. Bull. Math. Biol. 2007;69:1871–1886.
- [10] Korobeinikov A. Stability of ecosystem: global properties of a general prey-predator model. Math. Med. Biol. 2009;26:309–321.
- [11] Korobeinikov A. Global properties of a general predator-prey model with non-symmetric attack and consumption rate. Disc. Cont. Dyn. Syst. Ser. B. 2010;14:1095–1103.
- [12] McCluskey CC. Global stability for a class of mass action systems allowing for latency in tuberculosis. Math. Anal. Appl. 2008;338:518–535.
- [13] McCluskey CC. Complete global stability for an SIR epidemic model with delay distributed or discrete. Nonlinear Anal. B. 2010;11:55–59.
- [14] McCluskey CC. Global stability for an SEIR epidemiological model with varying infectivity and infinite delay. Math. Biosci. Eng. 2009;6:603–610.
- [15] Magal P, McCluskey CC, Webb GF. Liapunov functional and global asymptotic stability for an infection-age model. Appl. Anal. 2010;89:1109–1140.
- [16] Melnik AV, Korobeinikov A. Lyapunov functions and global stability for SIR and SEIR models with age-dependent susceptibility. Math. Biosci. Eng. 2013;10:369–378.
- [17] Nowak MA, May RM. Virus dynamics: mathematical principles of immunology and virology. New York (NY): Oxford University Press; 2000.
- [18] Shaikhet L. Lyapunov functionals and stability of stochastic functional differential equations. London: Springer; 2013.
- [19] Korobeinikov A. Global asymptotic properties of virus dynamics models with dose dependent parasite reproduction and virulence, and nonlinear incidence rate. Math. Med. Biol. 2009;26:225–239.
- [20] Beretta E, Kolmanovskii V, Shaikhet L. Stability of epidemic model with time delays influenced by stochastic perturbations. Math. Comput. Simul. 1998;45:269–277. Special Issue "Delay Systems".
- [21] Shaikhet L. Lyapunov functionals and stability of stochastic difference equations. London: Springer; 2011.
- [22] Caraballo T, Real J, Shaikhet L. Method of Lyapunov functionals construction in stability of delay evolution equations. J. Math. Anal. Appl. 2007;334:1130–1145.
- [23] Gantmacher FR. The theory of matrices. Vol. 2. New York (NY): Chelsea Pub. Co.; 1959.
- [24] Gikhman II, Skorokhod AV. Stochastic differential equations. Berlin: Springer; 1972.

#### Appendix 1.

Let us consider the Itô stochastic differential equation (see [24])

$$dx(t) = a_1(t, x(t))dt + a_2(t, x(t))dw(t),$$
  

$$t \ge 0, \quad x(t) \in \mathbf{R}^n, \quad x(0) = x_0.$$
(A1)

We assume that  $a_i(t, 0) \equiv 0, i = 1, 2$ , and hence Equation (A1) has the trivial solution. The generator

$$LV(t,x) = V_t(t,x) + \nabla V'(t,x)a_1(t,x) + \frac{1}{2}Tr\left[a'_2(t,x)\nabla^2 V(t,x)a_2(t,x)\right],$$
 (A2)

where

$$V_t = \frac{\partial u(t, x)}{\partial t}, \qquad \nabla V = \left(\frac{\partial V(t, x)}{\partial x_1}, \dots, \frac{\partial V(t, x)}{\partial x_n}\right)$$
$$\nabla^2 V = \left(\frac{\partial^2 V(t, x)}{\partial x_i \partial x_j}\right), \quad i, j = 1, \dots, n,$$

is associated with Equation (A1) (see [24]).

Definition A.1 The trivial solution of Equation (A1) is called:

- mean square stable, if for each  $\varepsilon > 0$  there exists  $\delta > 0$  such that  $\mathbf{E}|x(t, x_0)|^2 < \varepsilon$  holds for all  $t \ge 0$ , provided that  $\mathbf{E}|x_0|^2 < \delta$ ;
- asymptotically mean square stable, if it is mean square stable, and, for any initial value  $x_0$ ,  $\lim_{t\to\infty} \mathbf{E} |x(t, x_0)|^2 = 0$ ;
- stable in probability, if for any  $\varepsilon_1 > 0$  and  $\varepsilon_2 > 0$  there exists  $\delta > 0$  such that, for any initial value  $x_0$ , solution  $x(t, x_0)$  to Equation (A1) satisfies condition  $\mathbf{P}\{\sup_{t\geq 0} |x(t, x_0)| > \varepsilon_1\} < \varepsilon_2$ , where  $\mathbf{P}\{|x_0| < \delta\} = 1$ .

THEOREM A.1 Let there exist a function V(t, x) such that for any solution x(t) to Equation (A1) the following inequalities hold:

$$\mathbf{E}V(t, x(t)) \ge c_1 \mathbf{E}|x(t)|^2, \qquad t \ge 0,$$
(A1)

$$\mathbf{E}V(0, x_0) \le c_2 \mathbf{E}|x_0|^2,$$
 (A2)

$$\mathbf{E}LV(t, x(t)) \le -c_3 \mathbf{E}|x(t)|^2, \quad t \ge 0,$$
 (A3)

where  $c_i > 0$  and i = 1, 2, 3. Then the trivial solution of Equation (A1) is asymptotically mean square stable.

THEOREM A.2 Let there exist a function V(t, x) such that for any solution x(t) of the Equation (A1) the following inequalities hold:

$$V(t, x(t)) \ge c_1 |x(t)|^2,$$
 (A4)

$$V(0, x_0) \le c_2 |x_0|^2, \tag{A5}$$

$$LV(t, x(t)) < 0, \quad t > 0,$$
 (A6)

 $c_i > 0$ , i = 1, 2, for any initial value  $x_0$  such that  $\mathbf{P}\{|x_0| \le \delta\} = 1$ , where  $\delta > 0$  is small enough. Then the trivial solution of the equation of (A1) is stable in probability.

*Remark A.1* Let us assume, that for some nonlinear stochastic differential equation with an order of nonlinearity higher than one, there exists a function V(t, x), which satisfies Theorem A.1 hypotheses for the linear part (or the linear approximation) of the considered nonlinear equation. As it is shown in [18, p.131], this functional satisfies also Theorem A.2 hypotheses for the original nonlinear differential

equation. Thus, to get sufficient conditions for the stability in probability of the trivial solution of the nonlinear equation with an order of nonlinearity higher than one, it is enough, by the virtue of a function V(t, x) that satisfies Theorem A.1 hypotheses, to obtain sufficient conditions for the asymptotic mean square stability of the trivial solution of the linear part of the original nonlinear equation.

Consider the system of two stochastic differential equations without delays

$$\dot{x}_1(t) = a_{11}x_1(t) + a_{12}x_2(t) + \sigma_1 x_1(t)\dot{w}_1(t),$$
  
$$\dot{x}_2(t) = a_{21}x_1(t) + a_{22}x_2(t) + \sigma_2 x_2(t)\dot{w}_2(t),$$
 (A7)

where  $a_{ij}$ ,  $\sigma_i$ , i, j = 1, 2, are constants, and  $w_1(t)$ ,  $w_2(t)$  are mutually independent standard Wiener processes. Let

$$A = \begin{pmatrix} a_{11} & a_{12} \\ a_{21} & a_{22} \end{pmatrix}, \qquad \delta_i = \frac{1}{2}\sigma_i^2, \quad i = 1, 2,$$
  
$$Tr(A) = a_{11} + a_{22}, \qquad \det(A) = a_{11}a_{22} - a_{12}a_{21},$$
  
$$A_1 = \det(A) + a_{11}^2, \qquad A_2 = \det(A) + a_{22}^2.$$
(A8)

LEMMA A.1 Suppose that Tr(A) < 0, det(A) > 0,  $a_{12} \neq 0$  and

$$\delta_1 < \frac{|Tr(A)|\det(A)|}{A_2}, \qquad \delta_2 < \frac{|Tr(A)|\det(A) - A_2\delta_1|}{A_1 - |Tr(A)|\delta_1|}.$$
(A9)

Then the trivial solution of system (A7) is asymptotically mean square stable.

COROLLARY A.1 If  $a_{12} = 0$ , then conditions (A9) take the form

$$\delta_1 < -a_{11}, \quad \delta_2 < -a_{22}.$$

So, if the trivial solution of the first Equation (A7) is asymptotically mean square stable, then the stability condition of the trivial solution of the second Equation (A7) does not depend on  $a_{21}$ .