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Stochastic modeling of plant-insect interaction dynamics with MEMS-based monitoring and noise effects

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The dynamics of plant-insect interactions play a crucial role in the ecosystem, influenced by complex molecular signaling pathways. This study extends existing deterministic models of plant-insect systems by incorporating stochastic elements and molecular interactions, particularly focusing on the roles of Botrytis Induced Kinase-1 (BIK1) and Phyto Alexin Deficient-4 (PAD4) proteins. The model evaluates the effects of constant inhibition, pulsed inhibition, and adaptive feedback control on plant biomass (y_1), insect herbivore density (y_2), PAD4 levels (y_3), and BIK1 levels (y_4). Additionally, we examine the impact of different noise types, including deterministic, Gaussian, and Lévy noise, on system variability and stability. Results indicate that our stochastic model is superior as it shows a significant reduction in BIK1 levels, particularly under higher noise intensities, which enhances PAD4 activity and improves plant defense mechanisms. Moreover, moderate noise intensity ($\sigma = 0.05$) provides an optimal balance, sustaining PAD4 levels while effectively controlling insect herbivore populations. We also integrate MEMS-based feedback mechanisms, which dynamically adjust plant biomass and molecular signaling, further stabilizing the system's response to environmental variability.

KEYWORDS

mathematical analysis, feedback mechanism, immunity dynamics, Lévy noise, MEMS

1 Introduction

The intricate dynamics of plant-insect interactions have long captivated ecologists and biologists due to their profound implications for ecosystem stability and agricultural productivity. A notable example of such interaction is the relationship between plants and herbivorous insects, such as aphids, which are significant contributors to crop damage and yield loss globally [1]. Plants have evolved sophisticated defense mechanisms to counteract these insect attacks, initiating a complex interplay of molecular signals and physiological responses aimed at mitigating damage and ensuring survival [2, 3]. In this context, the role of Botrytis Induced Kinase-1 (BIK1) has emerged as a critical component of the plant defense arsenal. BIK1 is involved in the phosphorylation of the flagellin receptor FLS2 and BAK1 proteins, initiating a cascade of defense responses that include the production of phytoalexins and other defensive compounds [4]. These responses are modulated by signaling pathways mediated by jasmonic acid (JA) and salicylic acid (SA), which are crucial for the activation of plant immune responses [5, 6]. Recent research has highlighted the interaction between BIK1 and Phytoalexin Deficient-4 (PAD4), another

key player in the plant defense response. *PAD4* is known to enhance the plant's resistance to aphids by promoting the synthesis of antixenotic compounds that deter insect feeding [7]. However, *BIK1* has been shown to suppress *PAD4* expression, thereby modulating the plant's defensive capabilities [8]. Given the pivotal roles of *BIK1* and *PAD4* in plant-insect interactions, understanding the molecular underpinnings of their interaction and its impact on plant health is of paramount importance. We further enhance the deterministic model [9] by incorporating MEMS-based feedback mechanisms, allowing for real-time dynamic adjustments in plant biomass and molecular signaling based on environmental conditions and sensor data. Our study addresses critical gaps in existing research by incorporating stochastic elements to capture the inherent variability and extreme events in these systems, offering novel insights into how molecular signaling pathways influence plant defense mechanisms and herbivore population dynamics. The model simulates the dynamics of plant biomass (*P*), insect herbivore density (*I*), *PAD4* protein, and *BIK1* protein under various conditions. Mathematical modeling has evolved from deterministic approaches, such as those by [10, 11], to stochastic models that account for variability and randomness, as highlighted by [12, 13], and [14]. These stochastic models provide a more realistic depiction of ecological systems by incorporating environmental noise, as seen in works like [15, 16].

The stochastic processes, particularly Lévy noise, enhances the model's ability to capture extreme events, such as insect outbreaks, which are not well represented by Gaussian noise alone. Stochastic modeling techniques, as demonstrated by [17, 18], offer valuable insights for predicting population dynamics, evaluating control strategies, and understanding the influence of environmental variability. The use of Lévy noise in our model allows for the simulation of significant, abrupt changes in plant-insect interactions, providing a comprehensive representation of random phenomena. This approach underscores the importance of stochastic models in understanding complex biological systems and informing agricultural and environmental management practices. The characteristic function of a Lévy process is given by,

$$\mathbb{E} \left[e^{iuL(t)} \right] = \exp \left(t \left(iu\gamma - \frac{1}{2} \sigma^2 u^2 + \int_{\mathbb{R} \setminus \{0\}} (e^{iux} - 1 - iux \mathbf{1}_{|x| < 1}) \nu(dx) \right) \right),$$

where γ represents the drift coefficient, σ^2 is the variance of the Gaussian part, and ν is the Lévy measure that describes the jump intensity and distribution.

The plant-insect interaction system is a well-studied model in ecology, evolving from predator-prey analogies to more sophisticated mathematical models incorporating plant immunity concepts. This study extends these models by including molecular interactions in the plant defense system, inspired by [9]. The primary objective of this research is to develop a stochastic mathematical model to analyze plant-insect interaction dynamics, focusing on the molecular interplay between *PAD4* and *BIK1* proteins. It explores the impact of noise types, deterministic, Gaussian, and Lévy noise on the system's variability and stability. This research contributes to understanding plant-insect interactions at the molecular level, with potential applications in agriculture. It highlights the benefits of adaptive feedback control for plant protection, dynamically adjusting to changing conditions. By incorporating noise, especially Lévy noise, the model captures extreme fluctuations, providing a

realistic depiction of variability in plant-insect interactions. The study aims to answer key questions about the effects of control strategies, noise types, and the broader ecological implications of these findings.

2 Model equations

The system of differential equations governing the deterministic model is [9],

$$\begin{aligned} \frac{dy_1}{dt} &= a_1 y_1 \left(1 - \frac{y_1}{K} \right) - a_2 y_1 y_2, \\ \frac{dy_2}{dt} &= a_3 y_1 y_2 - a_4 y_2 - a_5 y_3 y_2, \\ \frac{dy_3}{dt} &= a_6 y_1 y_2 - a_7 y_3 - a_8 y_3 y_4, \\ \frac{dy_4}{dt} &= a_9 y_1 y_2 - a_{10} y_4 - a_{11} \cdot In_{BIK1}. \end{aligned} \quad (1)$$

The stochastic differential equations is,

$$\begin{aligned} dy_1 &= \left(a_1 y_1 \left(1 - \frac{y_1}{K} \right) - a_2 y_1 y_2 \right) dt + \sigma_1 y_1 dW_1, \\ dy_2 &= (a_3 y_1 y_2 - a_4 y_2 - a_5 y_3 y_2) dt + \sigma_2 y_2 dW_2, \\ dy_3 &= (a_6 y_1 y_2 - a_7 y_3 - a_8 y_3 y_4) dt + \sigma_3 y_3 dW_3, \\ dy_4 &= (a_9 y_1 y_2 - a_{10} y_4 - a_{11} \cdot In_{BIK1}) dt + \sigma_4 y_4 dW_4, \end{aligned} \quad (2)$$

The choice of noise addition is grounded in both biological and mathematical reasoning. Biologically, in ecological systems like plant-insect interactions, fluctuations in population densities and molecular levels are typically influenced by current population or protein levels [19–21]. Environmental stresses, such as insect outbreaks or weather conditions, do not affect the system uniformly but have a state-dependent effect: larger populations or protein levels experience more significant impacts. As a result, multiplicative noise (state-dependent noise) is biologically appropriate because it reflects that larger variables are more susceptible to noise. Mathematically, adding noise terms directly allows for stochastic perturbation while preserving the structure of the deterministic model. This approach simplifies the analysis and is commonly used in models involving stochastic dynamics through stochastic differential equations (SDEs), providing a meaningful representation of randomness.

PAD4 is a crucial gene that helps *Arabidopsis* plants defend against aphids. When aphids feed on the plant, *PAD4* gets activated, boosting the plant's defenses. *PAD4* helps in two main ways: by producing substances that deter aphids (antixenosis) and by creating chemicals that can harm aphids (antibiosis). *PAD4*'s activity is regulated by other genes like *TPS11* and *LOX5*. *TPS11* deals with sugar metabolism, while *LOX5* is involved in fatty acid metabolism. *TPS11*'s activity increases in the plant shoots when aphids attack, while *LOX5* activity increases in the leaves. Aphid attacks also trigger another gene, *MPL1*, which helps in defense but works independently of *PAD4*. The *BIK1* gene suppresses *PAD4*. If *BIK1* is less active (as in *BIK1* mutants), *PAD4* levels rise, making the plant more resistant to aphids. The interaction between *BIK1* and *PAD4* involves ethylene (ET), a plant hormone that also plays a role in defense. *PAD4* is essential for a full defense response against aphids, involving the production of deterrents and toxins. The plant's ability to emit ethylene is crucial for repelling aphids and

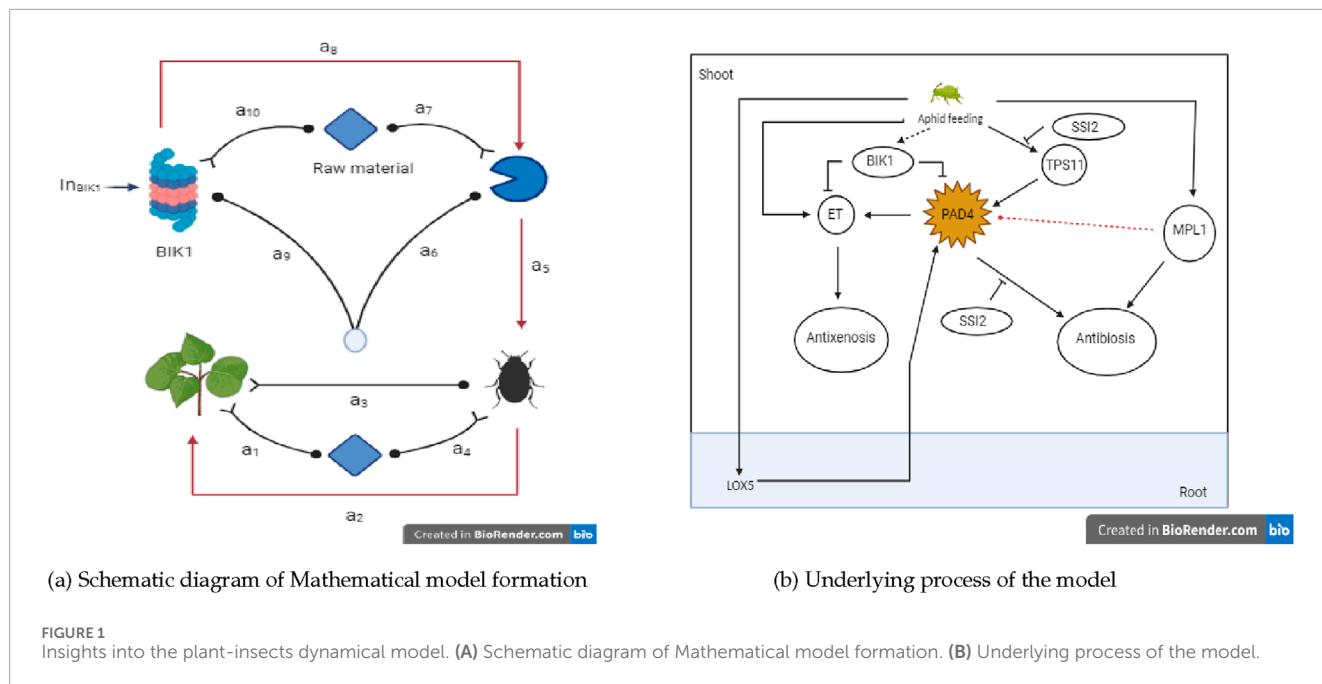


TABLE 1 Variables of the model.

S. No	System component	Description	Notation
1	P	Plant Biomass	γ_1
2	I	Insect herbivore density	γ_2
3	PAD4	Phytoalexin Deficient protein	γ_3
4	BIK1	Botrytis Induced Kinase protein	γ_4
5	In _{BIK1}	BIK1 inhibitor	In _{BIK1}

is dependent on PAD4 activity [22–27]. The mathematical model formation on this basis is shown in Figure 1A. This underlying process is shown in Figure 1B. The variables of the model are shown in Table 1.

The key assumptions of the model 1,2 are as follow,

- i. The model assumes that the plant biomass (P), insect herbivore density (I), PAD4 protein (PAD4), and BIK1 protein (BIK1) are homogeneously mixed within the environment. This means that these components are evenly distributed, and their interactions occur uniformly throughout the system.
- ii. The environmental conditions, such as temperature, humidity, and nutrient availability, are assumed to be constant.
- iii. The model considers a closed system with no immigration or emigration of insect herbivores. The population dynamics of the insect herbivores are governed solely by the birth and death rates within the system.
- iv. The concentration of the BIK1 inhibitor (In_{BIK1}) is assumed to be constant and does not change over time.
- v. The rate constants (a_i) and carrying capacity (K) are assumed to be constant over time.

- vi. The model does not consider time delays in the responses of the components. All interactions and changes occur instantaneously.
- vii. The model assumes that there are no external interventions, such as pesticide applications or genetic modifications.

3 Qualitative analysis of the model

3.1 Existence and uniqueness of the solution

The following section provides existence, boundedness, and equilibrium analysis for the current model. Theorem 3.1 ensures that the interactions between variables of the system are well-defined and consistent under all modeled conditions, making the system biologically predictable. While the approach used here is specific to our system, similar results in other contexts can be found in [28, 29].

TABLE 2 Parameter values used in the model.

Description	Rate constants	Values of rate constants	References
Plant biomass production rate	a_1	0.2	[30, 31]
Insect infestation on plants	a_2	0.6	[30, 31]
Insect reproduction rate	a_3	0.01	[30, 31]
Insect death rate	a_4	0.02	[30, 31]
Antixenosis by <i>PAD4</i>	a_5	0.002	[32]
<i>PAD4</i> production	a_6	1	Assumed
<i>PAD4</i> degradation	a_7	0.1	[33, 34]
<i>BIK1</i> mediated <i>PAD4</i> decrease	a_8	0.1	[35]
<i>BIK1</i> production	a_9	1	[36–38]
<i>BIK1</i> degradation	a_{10}	0.1	Assumed
Inhibitor based <i>BIK1</i> decrease	a_{11}	0.1	Assumed
Carrying capacity of plant biomass	K	1.0	Assumed
Noise intensity	σ	0.01	Assumed

Theorem 3.1: For system 1, there exists a unique solution.

Proof. The proof is given in appendix section (Theorem 3.1).

Theorem 3.2: The solutions of the system 1 are bounded for all $t \geq 0$.

Proof. The proof is given in appendix section (Theorem 3.1). Theorem 3.2 reflects that the population levels of plants, insects, and proteins will not grow indefinitely or collapse to zero, showing the natural limits on growth due to environmental factors and resource constraints. The proof is given in appendix section (Theorem 3.2).

3.2 Equilibrium analysis

In this section, we analyze the equilibrium points of the system and assess their stability. The variational matrix is a matrix of first-order partial derivatives that encapsulates the local linearization of a nonlinear dynamical system around its equilibrium points. If the system is described by a set of ordinary differential equations $\dot{\mathbf{y}} = f(\mathbf{y})$, where $\mathbf{y} = (y_1, y_2, \dots, y_n)$ represents the state variables, the variational matrix V is given by,

$$V = \frac{\partial f(\mathbf{y})}{\partial \mathbf{y}} = \begin{pmatrix} \frac{\partial f_1}{\partial y_1} & \frac{\partial f_1}{\partial y_2} & \dots & \frac{\partial f_1}{\partial y_n} \\ \frac{\partial f_2}{\partial y_1} & \frac{\partial f_2}{\partial y_2} & \dots & \frac{\partial f_2}{\partial y_n} \\ \vdots & \vdots & \ddots & \vdots \\ \frac{\partial f_n}{\partial y_1} & \frac{\partial f_n}{\partial y_2} & \dots & \frac{\partial f_n}{\partial y_n} \end{pmatrix}.$$

This matrix captures the infinitesimal behavior of the system around the equilibrium points by linearizing the system. The

eigenvalue spectrum of the variational matrix governs the local stability properties of the equilibrium. If all eigenvalues have negative real parts, the system exhibits asymptotic stability at the equilibrium point. If any eigenvalue has a positive real part, the equilibrium is unstable. The variational matrix of our model is given by,

$$V = \begin{pmatrix} a_1 \left(1 - \frac{2}{K}\right) - a_2 & -a_2 & 0 & 0 \\ a_3 & a_3 - a_4 - a_5 & -a_5 & 0 \\ a_6 & a_6 & -a_7 - a_8 & -a_8 \\ a_9 & a_9 & 0 & -a_{10} \end{pmatrix}.$$

3.2.1 $\mathcal{E}_1(0, 0, 0, 0)$

The variational matrix V at $\mathcal{E}_1(0, 0, 0, 0)$ is,

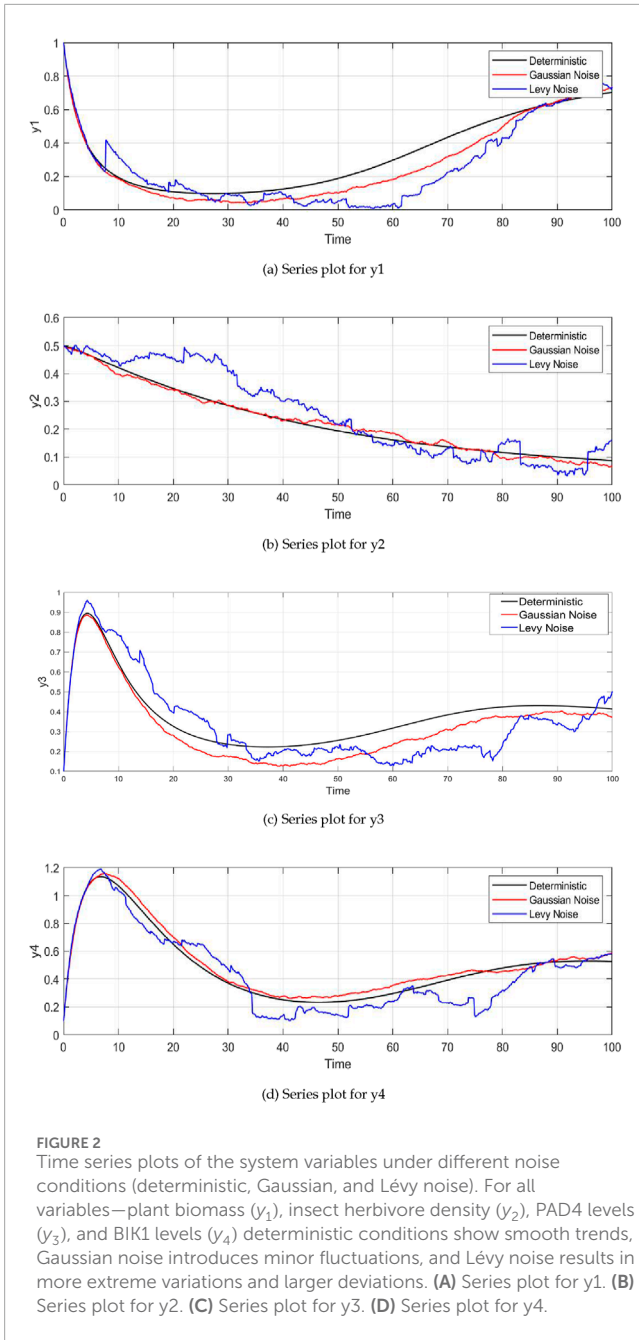
$$V|_{\mathcal{E}_1(0,0,0,0)} = \begin{pmatrix} a_1 & 0 & 0 & 0 \\ 0 & -a_4 & 0 & 0 \\ 0 & 0 & -a_7 & 0 \\ 0 & 0 & 0 & -a_{10} \end{pmatrix}.$$

So, the eigenvalues at the equilibrium point $\mathcal{E}_1(0, 0, 0, 0)$ are,

$$\lambda_1 = a_1, \quad \lambda_2 = -a_4, \quad \lambda_3 = -a_7, \quad \lambda_4 = -a_{10}$$

Theorem 3.3: The equilibrium point $\mathcal{E}_1(0, 0, 0, 0)$ is locally asymptotically stable, if $a_1 < 0$.

Proof. The proof is easy to follow.



3.2.2 $E_2(1,0,0,0)$

The variational matrix V at $E_2(1,0,0,0)$ is,

$$V|_{E_2(1,0,0,0)} = \begin{pmatrix} a_1 \left(1 - \frac{2}{K}\right) & -a_2 & 0 & 0 \\ 0 & a_3 - a_4 & 0 & 0 \\ 0 & a_6 & -a_7 & 0 \\ 0 & a_9 & 0 & -a_{10} \end{pmatrix}.$$

So, the eigenvalues at the equilibrium point $E_2(1,0,0,0)$ are,

$$\lambda_1 = a_1 \left(1 - \frac{2}{K}\right), \quad \lambda_2 = a_3 - a_4, \quad \lambda_3 = -a_7, \quad \lambda_4 = -a_{10}$$

Theorem 3.4: The equilibrium point $E_2(1,0,0,0)$ is locally asymptotically stable, if $K < 2$ and $a_1 < a_2$ or $a_3 < a_4 + a_5$.

Proof. The proof is easy to follow.

3.2.3 $E_3(0,1,0,0)$

The variational matrix for $E_3(0,1,0,0)$ is,

$$V|_{E_3(0,1,0,0)} = \begin{pmatrix} a_1 - a_2 & 0 & 0 & 0 \\ a_3 & -a_4 & -a_5 & 0 \\ a_6 & 0 & -a_7 & 0 \\ a_9 & 0 & 0 & -a_{10} \end{pmatrix}.$$

The eigenvalues are for the equilibrium point $E_3(0,1,0,0)$ are,

$$\lambda_1 = a_1 - a_2, \quad \lambda_2 = -a_4, \quad \lambda_3 = -a_7, \quad \lambda_4 = -a_{10}$$

Theorem 3.5: The equilibrium point $E_3(0,1,0,0)$ is locally asymptotically stable, if $a_1 < a_2$.

Proof. The proof is easy to follow.

3.2.4 $E_4(0,0,1,0)$

The variational matrix for $E_4(0,0,1,0)$ is,

$$V|_{E_4(0,0,1,0)} = \begin{pmatrix} a_1 & 0 & 0 & 0 \\ 0 & -a_4 - a_5 & 0 & 0 \\ 0 & 0 & -a_7 & -a_8 \\ 0 & 0 & 0 & -a_{10} \end{pmatrix}.$$

The eigenvalues are for the equilibrium point $E_4(0,0,1,0)$ are,

$$\lambda_1 = a_1, \quad \lambda_2 = -a_4 - a_5, \quad \lambda_3 = -a_7, \quad \lambda_4 = -a_{10}$$

Theorem 3.6: The equilibrium point $E_4(0,0,1,0)$ is locally asymptotically stable, if $a_1 < 0$.

Proof. The proof is easy to follow.

3.2.5 $E_5(0,0,0,1)$

The variational matrix for $E_5(0,0,0,1)$ is,

$$V|_{E_5(0,0,0,1)} = \begin{pmatrix} a_1 & 0 & 0 & 0 \\ 0 & -a_4 & 0 & 0 \\ 0 & 0 & -a_7 - a_8 & 0 \\ 0 & 0 & 0 & -a_{10} \end{pmatrix}.$$

The eigenvalues are for the equilibrium point $E_5(0,0,0,1)$ are,

$$\lambda_1 = a_1, \quad \lambda_2 = -a_4, \quad \lambda_3 = -a_7 - a_8, \quad \lambda_4 = -a_{10}$$

Theorem 3.7: The equilibrium point $E_5(0,0,0,1)$ is locally asymptotically stable, if $a_1 < 0$.

Proof. The proof is easy to follow.

3.2.6 $E_6(1,1,0,0)$

The variational matrix for $E_6(1,1,0,0)$ is,

$$V|_{E_6(1,1,0,0)} = \begin{pmatrix} a_1 \left(1 - \frac{2}{K}\right) - a_2 & -a_2 & 0 & 0 \\ a_3 & a_3 - a_4 & 0 & 0 \\ a_6 & a_6 & -a_7 & 0 \\ a_9 & a_9 & 0 & -a_{10} \end{pmatrix}.$$

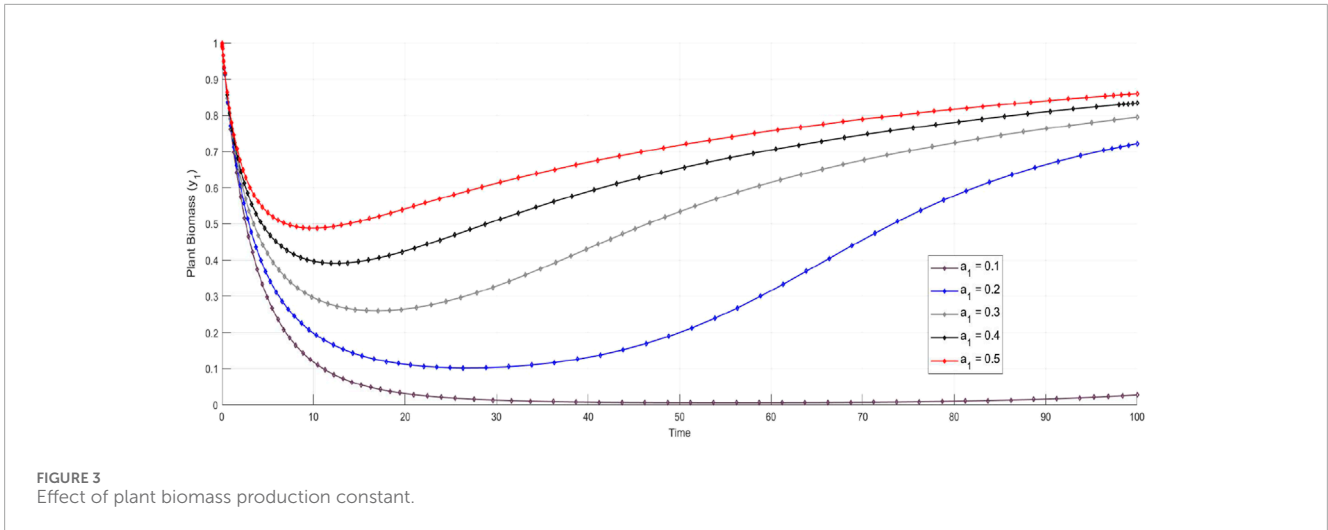


FIGURE 3 Effect of plant biomass production constant.

The eigenvalues are for the equilibrium point $\mathcal{E}_6(1, 1, 0, 0)$ are,

$$\lambda_1 = a_1 \left(1 - \frac{2}{K}\right) - a_2, \quad \lambda_2 = a_3 - a_4, \quad \lambda_3 = -a_7, \quad \lambda_4 = -a_{10}$$

Theorem 3.8: The equilibrium point $\mathcal{E}_6(1, 1, 0, 0)$ is locally asymptotically stable, if $a_1 \left(1 - \frac{2}{K}\right) < a_2$ and $a_3 < a_4$.

Proof. The proof is easy to follow.

3.2.7 $\mathcal{E}_7(1, 0, 1, 0)$

The variational matrix at $\mathcal{E}_7(1, 0, 1, 0)$ is,

$$V|_{\mathcal{E}_7(1,0,1,0)} = \begin{pmatrix} a_1 \left(1 - \frac{2}{K}\right) & 0 & 0 & 0 \\ 0 & -a_4 - a_5 & 0 & 0 \\ 0 & a_6 & -a_7 & 0 \\ 0 & 0 & 0 & -a_{10} \end{pmatrix}.$$

The eigenvalues are for the equilibrium point $\mathcal{E}_7(1, 0, 1, 0)$ are,

$$\lambda_1 = a_1 \left(1 - \frac{2}{K}\right), \quad \lambda_2 = -a_4 - a_5, \quad \lambda_3 = -a_7, \quad \lambda_4 = -a_{10}$$

Theorem 3.9: The equilibrium point $\mathcal{E}_7(1, 0, 1, 0)$ is locally asymptotically stable, if $1 - \frac{2}{K} < 0$ (i.e., $K < 2$).

Proof. The proof is easy to follow.

3.2.8 $\mathcal{E}_8(1, 0, 1, 1)$

The variational matrix at $\mathcal{E}_8(1, 0, 1, 1)$ is,

$$V|_{\mathcal{E}_8(1,0,1,1)} = \begin{pmatrix} a_1 \left(1 - \frac{2}{K}\right) & 0 & 0 & 0 \\ 0 & -a_4 - a_5 & 0 & 0 \\ 0 & a_6 & -a_7 - a_8 & -a_8 \\ 0 & 0 & 0 & -a_{10} \end{pmatrix}.$$

The eigenvalues are for the equilibrium point $\mathcal{E}_8(1, 0, 1, 1)$ are,

$$\lambda_1 = a_1 \left(1 - \frac{2}{K}\right), \quad \lambda_2 = -a_4 - a_5, \quad \lambda_3 = -a_7 - a_8, \quad \lambda_4 = -a_{10}$$

Theorem 3.10: The equilibrium point $\mathcal{E}_8(1, 0, 1, 1)$ is locally asymptotically stable, if $1 - \frac{2}{K} < 0$ (i.e., $K < 2$).

Proof. The proof is easy to follow.

3.3 Biological significance

The equilibrium analysis emphasizes the essential dynamics between plant biomass, insect herbivores, and defense proteins, PAD4 and BIK1, in maintaining ecological stability. Theorems 3.4, 3.5 establish the foundational roles of plants and insects in the system, highlighting how their coexistence is necessary for sustaining balanced populations. The most biologically significant results are demonstrated in Theorems 3.9, 3.10, where plants maintain stable coexistence with PAD4 or both PAD4 and BIK1 proteins. These equilibria reflect the plant's defensive mechanisms being actively regulated by these proteins, ensuring preparedness for potential herbivore attacks. Theorem 3.8 presents the ecologically balanced state, where both plants and insect herbivores coexist, with PAD4 and BIK1 proteins playing a regulatory role. This equilibrium ensures that plant defense systems, driven by these proteins, manage herbivore populations effectively, maintaining system stability. This section highlights the crucial role of plant defense proteins in regulating insect interactions, ensuring long-term ecological balance.

3.4 Basic reproduction number

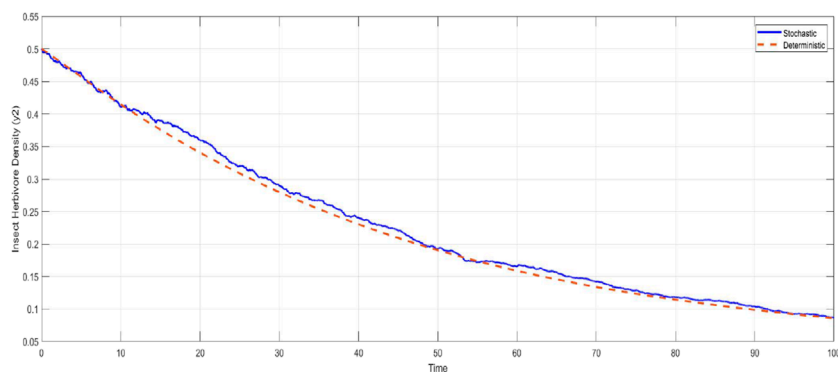
The basic reproduction number R_0 is given by,

$$R_0 = \frac{a_3 K}{a_4}.$$

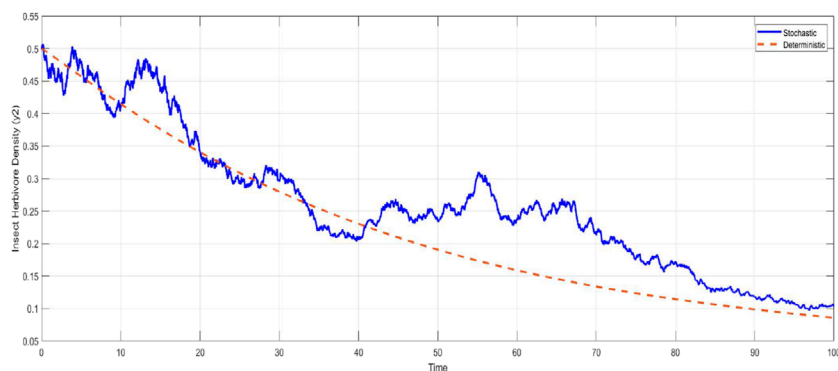
Theorem 3.11: If $R_0 > 1$, then the system is uniformly persistent, meaning there exists a positive constant δ such that for any solution $(y_1(t), y_2(t), y_3(t), y_4(t))$ with initial conditions in the interior of the positive orthant, we have,

Proof. The proof is given in appendix (Theorem 3.11).

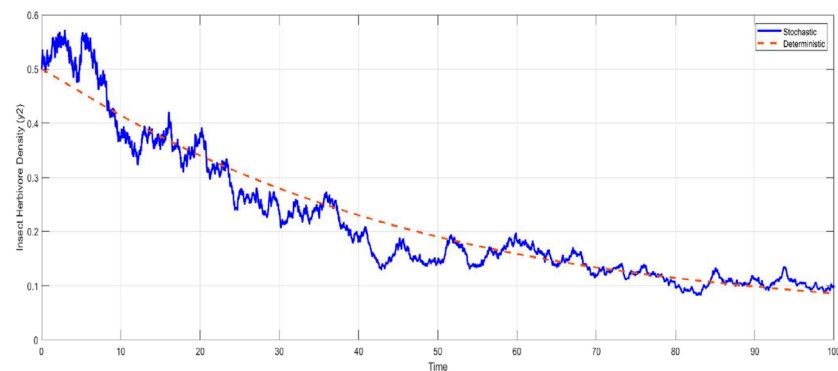
$$\liminf_{t \rightarrow \infty} y_i(t) \geq \delta > 0, \quad \text{for } i = 1, 2, 3, 4.$$



(a) Series plot for $\sigma=0.01$



(b) Series plot for $\sigma=0.05$



(c) Series plot for $\sigma=0.1$

FIGURE 4 Time series of insect herbivore density (y_2) under different noise intensities. With higher noise intensity ($\sigma = 0.1$), the fluctuations are significantly larger, leading to increased variability in the system's behavior. **(A)** Series plot for $\sigma=0.01$. **(B)** Series plot for $\sigma=0.05$. **(C)** Series plot for $\sigma=0.1$.

3.5 Stochastic analysis

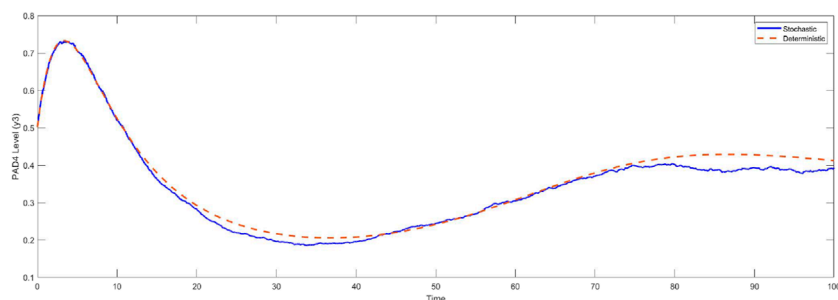
For stochastic system with y_1, y_2, y_3, y_4 , we define Γ as

$$\Gamma = \left\{ (y_1(t), y_2(t), y_3(t), y_4(t)) \in \mathbb{R}_+^4 : y_1(t) + y_2(t) + y_3(t) + y_4(t) \leq \frac{a_1}{a_2} \right\}$$

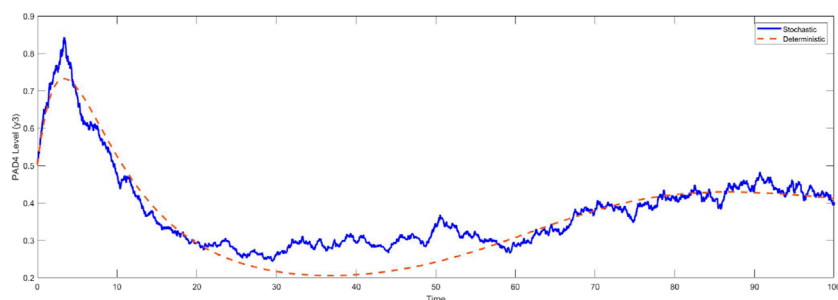
We need to show that Γ fulfills almost sure invariance principle.

Theorem 3.12: *The closed set Γ fulfills almost sure invariance principle for the stochastic system 2.*

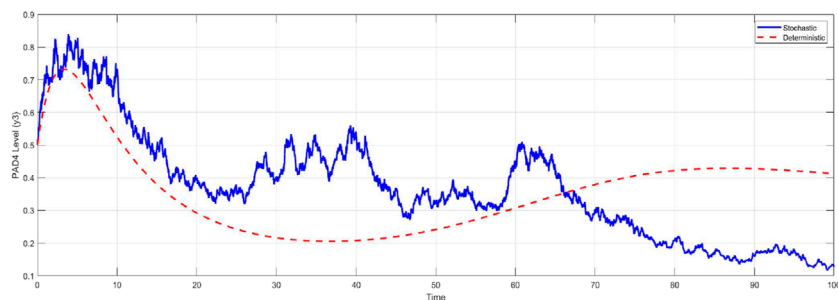
Proof. This theorem guarantees that the system remains within realistic bounds even under stochastic influences, indicating that the



(a) Series plot for $\sigma=0.01$



(b) Series plot for $\sigma=0.05$



(c) Series plot for $\sigma=0.1$

FIGURE 5

Time series of PAD4 protein levels (y_3) under different noise intensities. With higher noise intensity ($\sigma = 0.1$), significant fluctuations occur, showing increased sensitivity to the noise. (A) Series plot for $\sigma=0.01$. (B) Series plot for $\sigma=0.05$. (C) Series plot for $\sigma=0.1$.

ecosystem is robust to fluctuations in population and protein levels. The proof is given in appendix section (Theorem 3.12).

Theorem 3.13: For $(y_1(0), y_2(0), y_3(0), y_4(0)) \in \Gamma$, system 2 has a unique and positive solution almost surely.

Proof. This theorem assures that despite randomness, the system's biological variables (plants, insects, proteins) maintain positive values, ensuring the ecological system remains functional. The proof is given in appendix section (Theorem 3.13).

4 Numerical simulations

In the following subsections, various numerical results are provided. The values of the parameters used in the model

are given in Table 2. The Euler-Maruyama method is used to solve the stochastic differential equations iteratively as follow,

$$\begin{aligned}
 y_1(t + \Delta t) &= y_1(t) + \left(k_1 y_1(t) \left(1 - \frac{y_1(t)}{K} \right) - k_2 y_1(t) y_2(t) \right) \Delta t + \sigma \sqrt{\Delta t} \cdot \eta_1(t), \\
 y_2(t + \Delta t) &= y_2(t) + (k_3 y_1(t) y_2(t) - k_4 y_2(t) - k_5 y_3(t) y_2(t)) \Delta t + \sigma \sqrt{\Delta t} \cdot \eta_2(t), \\
 y_3(t + \Delta t) &= y_3(t) + (k_6 y_1(t) y_2(t) - k_7 y_3(t) - k_8 y_3(t) y_4(t)) \Delta t + \sigma \sqrt{\Delta t} \cdot \eta_3(t), \\
 y_4(t + \Delta t) &= y_4(t) + (k_9 y_1(t) y_2(t) - k_{10} y_4(t) - k_{11} \cdot 0.1) \Delta t + \sigma \sqrt{\Delta t} \cdot \eta_4(t),
 \end{aligned}$$

where $\eta_i(t)$ represents the noise term. For Gaussian noise, $\eta_i(t)$ is from a normal distribution $\mathcal{N}(0, 1)$,

$$\eta_i(t) \sim \mathcal{N}(0, 1).$$

For Lévy noise, $\eta_i(t)$ is from a Lévy distribution.

$$\eta_i(t) \sim \text{Lvy}(\alpha, \beta).$$

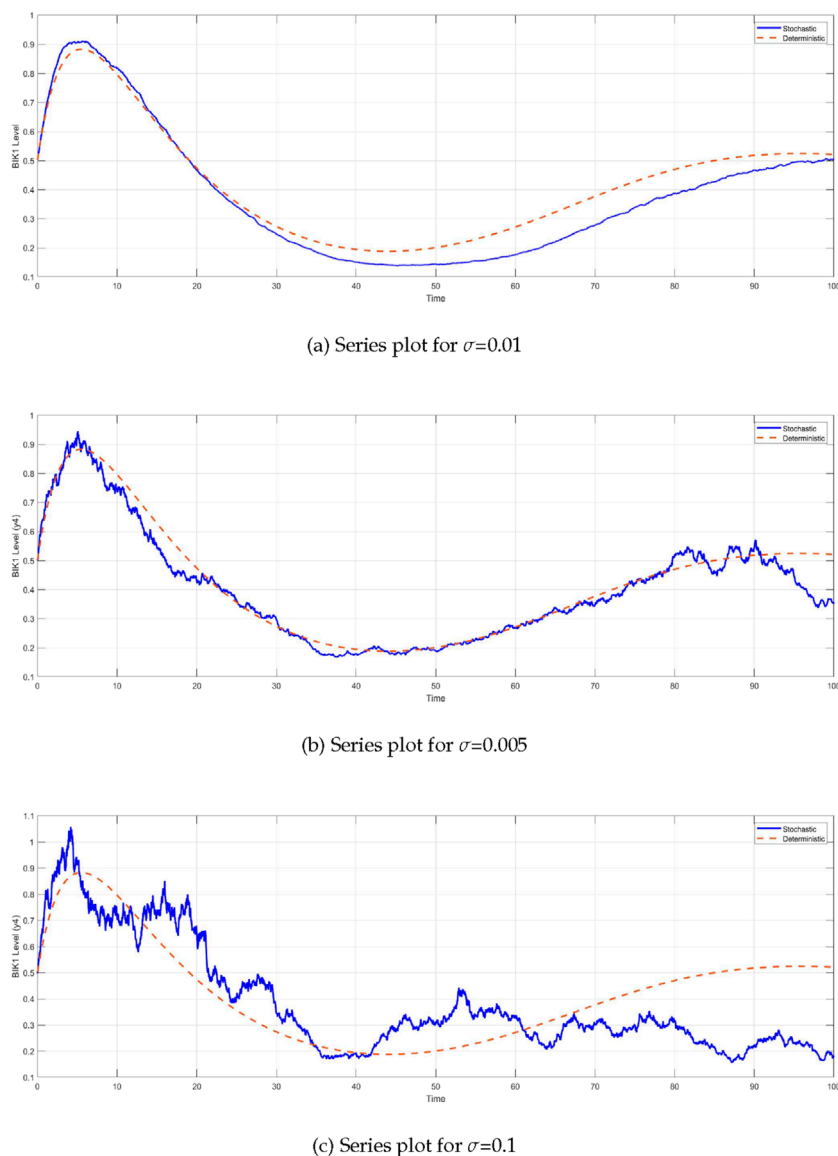


FIGURE 6 Time series of BIK1 protein levels (y_4) under different noise intensities. At higher noise intensity ($\sigma = 0.1$), the system exhibits large variability, indicating a stronger response to the perturbations. **(A)** Relationship between PAD4 Levels (y_3) and BIK1 Levels (y_4). **(B)** Relationship between Plant Biomass (y_1) and Insect Herbivore Density (y_2).

where,

$$\Delta W_t \sim \mathcal{N}(0, \Delta t).$$

And

$$\Delta W_t = \sqrt{\Delta t} \cdot Z,$$

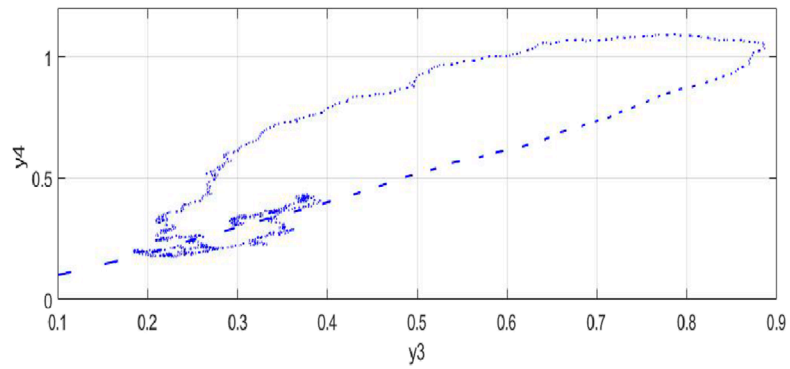
where Z is a standard normal random variable ($Z \sim \mathcal{N}(0, 1)$).

4.1 Noises comparison

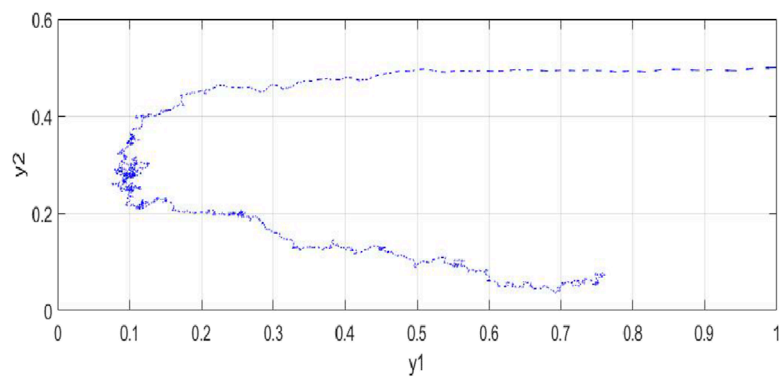
Figure 2A shows the dynamics of plant biomass (y_1) under deterministic, Gaussian, and Lévy noise conditions. Under

deterministic conditions, plant biomass follows a smooth decline and recovery trajectory. When Gaussian noise is introduced, the system exhibits more fluctuations compared to the deterministic case but maintains a generally similar trend. Lévy noise, however, results in significantly larger fluctuations, reflecting more extreme variations in plant biomass. This observation is consistent with findings from the study [39], which highlighted the importance of Lévy noise in capturing extreme events in biological systems.

The dynamics of insect herbivore density (y_2) under different noise conditions are illustrated in Figure 2B. Deterministic conditions show a continuous decline in insect density. Gaussian noise introduces more variability into the system, causing minor fluctuations around the declining trend. Lévy noise, on the other hand, introduces significant variability, leading to more pronounced



(a) Relationship between *PAD4* Levels (y_3) and *BIK1* Levels (y_4)



(b) Relationship between Plant Biomass (y_1) and Insect Herbivore Density (y_2)

FIGURE 7

The plot shows the relationships in the plant-insect interaction model: **(A)** between *PAD4* Levels (y_3) and *BIK1* Levels (y_4), and **(B)** between Plant Biomass (y_1) and Insect Herbivore Density (y_2).

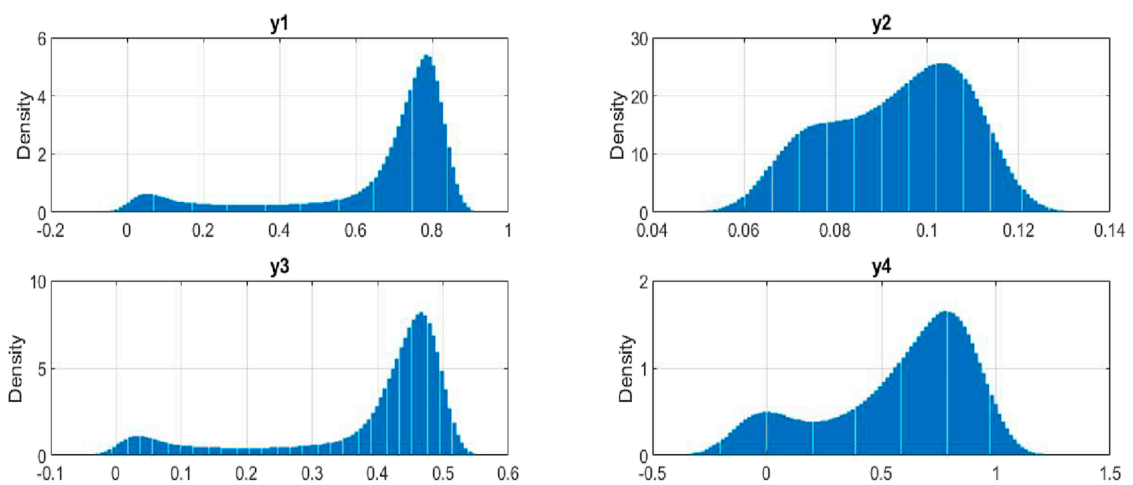
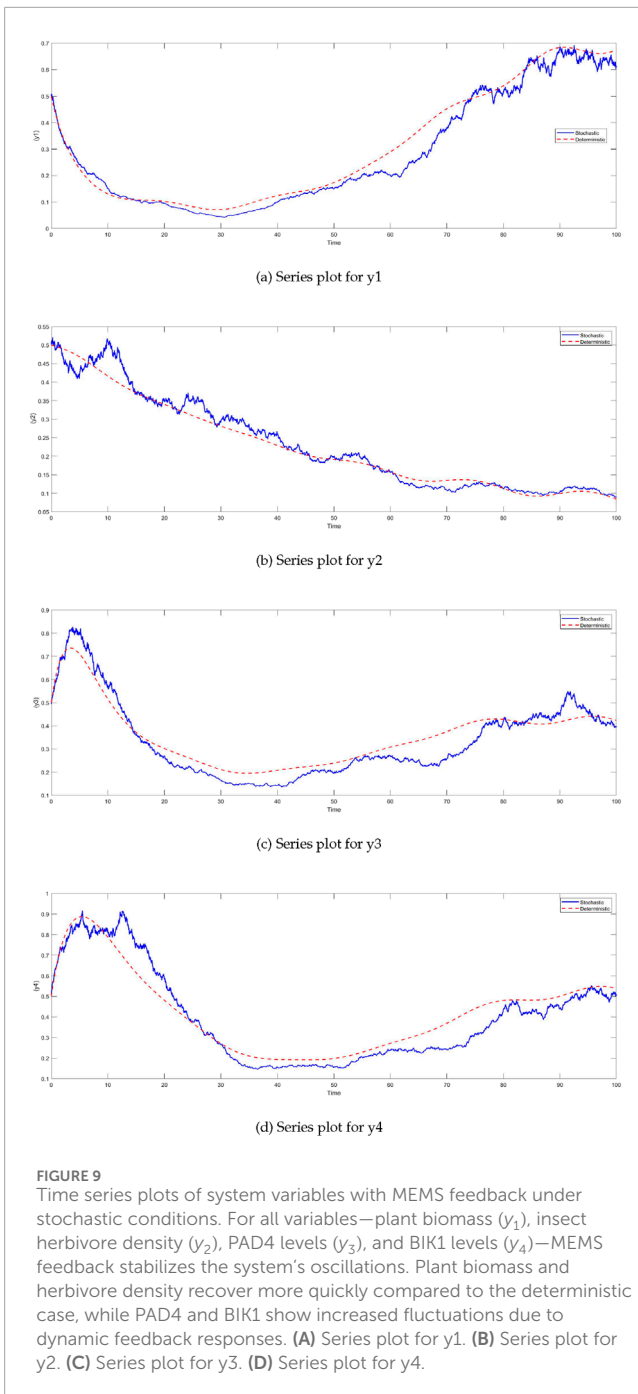


FIGURE 8

Density Distribution of the model variable's.



fluctuations in insect density. The same trend is followed by other variables (Figures 2C, D).

The parameter a_1 significantly impacts plant biomass y_1 (Figure 3A). Higher values of a_1 indicate better soil conditions and other suitable conditions, allowing plants to maintain or increase their biomass over time.

4.2 Lévy noise

The insect herbivore density (y_2) exhibits distinct behaviors under different noise intensities (Figure 4). At low noise

intensity ($\sigma = 0.01$) (Figure 4A), the herbivore density decreases steadily, closely mirroring the deterministic model. However, as noise intensity increases, particularly at moderate levels ($\sigma = 0.05$) (Figure 4B), there is an observable increase in fluctuations around the declining trend.

Similarly, the PAD4 levels (y_3) show an interesting pattern under varying noise intensities (Figure 5). At a moderate noise intensity of $\sigma = 0.05$ (Figure 5A), PAD4 levels maintain a relatively stable and higher average compared to both lower and higher noise intensities. This stability at moderate noise levels suggests that the system can better sustain its defense mechanisms, making this noise intensity particularly beneficial for maintaining PAD4 activity. Conversely, at higher noise intensities ($\sigma = 0.1$) (Figure 5C), PAD4 levels decrease significantly, indicating that excessive noise can disrupt the plant's defense responses.

The comparison between the deterministic and stochastic models reveals critical insights into the behavior of the system under different noise intensities (Figure 6). One of the most notable observations is the impact of stochasticity on the BIK1 levels (y_4). As noise intensity increases, BIK1 levels exhibit significant fluctuations. For higher noise intensities, BIK1 levels decrease more rapidly, indicating that the stochastic model is more sensitive to perturbations. This sensitivity suggests that in real-world scenarios, BIK1 levels are likely to be more variable due to environmental and internal noise, which the deterministic model fails to capture (Figures 6A–C) [40, 41].

4.3 Queir plots

Queir plots visually represent how the states of a system evolve over time, especially in complex, nonlinear systems. These plots show the system's trajectory or path in a simplified, multi-dimensional space, focusing on key variables.

Theorem 4.1: Consider the system defined by,

$$\frac{dy_1}{dt} = a_1 y_1 \left(1 - \frac{y_1}{K}\right) - a_2 y_1 y_2.$$

If $y_2(t) \leq \frac{a_1}{a_2} \left(1 - \frac{y_1}{K}\right)$, the plant biomass $y_1(t)$ will grow.

Proof. The proof is given in appendix section (Theorem 4.1).

Theorem 4.2: Consider the system defined by,

$$\frac{dy_3}{dt} = a_6 y_1 y_2 - a_7 y_3 - a_8 y_3 y_4,$$

$$\frac{dy_4}{dt} = a_9 y_1 y_2 - a_{10} y_4 - a_{11} \cdot \ln_{BIK1}.$$

If $y_1 y_2$ exceeds the thresholds $\frac{a_7 y_3 + a_8 y_3 y_4}{a_6}$ and $\frac{a_{10} y_4 + a_{11} \cdot \ln_{BIK1}}{a_9}$, both $y_3(t)$ and $y_4(t)$ will increase, contributing to the plant's defensive response.

Proof. The proof is given in appendix section (Theorem 4.2).

4.3.1 Relationship between PAD4 levels (y_3) and BIK1 levels (y_4)

As shown in Figure 7A, the relationship between PAD4 (y_3) and BIK1 (y_4) levels exhibits a complex trend where increases in PAD4 lead to increases in BIK1 up to a threshold, after

which the relationship stabilizes. This pattern reflects regulatory feedback mechanisms, where *PAD4* activation upregulates *BIK1* until feedback stabilization occurs. The findings align with studies on plant signaling pathways, where *BIK1* is essential in defense responses, and stochastic dynamics impact regulatory networks.

4.3.2 Relationship between plant biomass (y_1) and insect herbivore density (y_2)

Figure 7B shows that the relationship between plant biomass (y_1) and insect herbivore density (y_2) is inversely correlated, where increases in plant biomass lead to reductions in herbivore density. This reflects the plant's defense mechanisms, mediated by proteins like *PAD4* and *BIK1*, which regulate herbivore populations. Stochastic fluctuations observed are consistent with known plant-herbivore interaction models.

4.4 Probability density distributions

The stationary distribution describes the long-term behavior of a stochastic process, offering insights into the stability of system. A Markov process, which models a sequence of potential events, is primarily influenced by the state attained in the previous event. In simpler terms, it can be thought of as “what happens in the future depends only on the current situation.”

In the space \mathbb{R}_+^m , consider that the process $\mathbf{J}(t)$ is regular and time-homogeneous, exhibiting Markovian behavior with respect to time t of the form:

$$d\mathbf{J}(t) = \sum_{s=1}^l \kappa_s d\mathcal{B}_s(t) + b(\mathbf{J}) dt.$$

Here, $\mathcal{A}(\mathbf{J}) = [\iota_{ij}(y)]$ is the matrix associated with the mixing terms, where $\iota_{ij}(y) = \sum_{s=1}^l \kappa_s^i(y) \kappa_s^j(y)$.

Lemma 4.1: [42, 43] *The process $\mathbf{J}(t)$ is said to possess a unique stationary distribution $m(\cdot)$ if we can identify a bounded domain with regular boundaries $U, \bar{U} \subset \mathbb{R}^d$, such that,*

1. *The smallest eigenvalue of $\mathcal{A}(t)$ is bounded away from zero within the domain U and its vicinity.*

Additionally, if $y \in \mathbb{R}^d \setminus U$, the mean time τ (at which a path originating from y reaches U) is finite, and $\sup_{y \in L} \mathbb{E}\tau < \infty$ for each compact subset $L \subset \mathbb{R}^m$. Furthermore, let $g(\cdot)$ be an integrable function with respect to the measure π , then

$$\lim_{T \rightarrow \infty} \frac{1}{T} \int_0^T f(\mathbf{J}_y(t)) dt = \int_{\mathbb{R}^d} g(y) \pi(dy) = 1 \quad \text{for all } y \in \mathbb{R}^d.$$

we define,

$$\mathcal{R}_0^p = \frac{a_1 a_3 a_6}{\left(a_2 + \frac{c_1^2}{2}\right) \left(a_4 + \frac{c_2^2}{2}\right) \left(a_7 + \frac{c_3^2}{2}\right) \left(a_{10} + \frac{c_4^2}{2}\right)}. \quad (3)$$

Theorem 4.3: *For $(y_1(0), y_2(0), y_3(0), y_4(0)) \in \Gamma$, the system has a unique stationary distribution $\pi(\cdot)$ as well as the solution $(y_1(t), y_2(t), y_3(t), y_4(t))$ to the model is ergodic in nature.*

Proof. The proof is given in appendix section (Theorem 4.3).

Figure 8 presents the probability density distributions for the key variables in our stochastic model of plant-insect interactions. The

density of y_1 (is predominantly concentrated around 0.5, indicating that this is the most common biomass level. y_2 exhibits its highest density at approximately 0.1, showing that insect populations reach this density influenced by factors such as available plant biomass and natural predator presence. For y_3 , the density peaks sharply, demonstrating that *PAD4* protein levels consistently reach an optimal level for effective response to insect herbivory. In contrast, y_4 displays a bimodal distribution, indicating two dominant states of this protein. The lower levels of *BIK1*, corresponding to the first peak, are associated with heightened *PAD4* activity, reflecting a robust plant defense mechanism. The second peak at higher *BIK1* levels suggests a regulatory balance where *BIK1* moderates the defense response to prevent overreaction.

5 MEMS in control

The integration of Micro-Electromechanical Systems (MEMS) in plant systems has revolutionized the field of precision agriculture by providing real-time monitoring and adaptive control capabilities. MEMS sensors are widely used for tracking environmental parameters such as soil moisture, temperature, humidity, and nutrient levels, enabling more efficient and precise irrigation and fertilization strategies. For example, The increasing demand for the miniaturization of biosensors has driven growing interest in microelectromechanical systems (MEMS) [44, 45], along with nanoelectromechanical systems (NEMS) and microfluidic or lab-on-a-chip based biosensors [35, 46]. These compact systems provide enhanced accuracy, sensitivity, specificity, and cost-efficiency, while also offering high-performance biosensing capabilities. MEMS-based biosensors leverage a range of detection methods, including optical, mechanical, magnetic, and electrochemical approaches. For optical detection, probes like organic dyes, semiconductor quantum dots, and other fluorescence markers are commonly employed. In magnetic MEMS biosensors, nanoparticles such as magnetic, paramagnetic, or ferromagnetic particles are utilized. Mechanical MEMS biosensors operate based on changes in surface stress or mass [47], where biochemical reactions or analyte adsorption on the cantilever induce surface stress changes. Electrochemical MEMS biosensors, on the other hand, rely on amperometric, potentiometric, or conductometric detection methods [48–51]. For each variable, sensor-based feedback terms can be introduced to adjust the differential equations based on real-time data gathered by MEMS. This can include intervention strategies or feedback loops that modify plant biomass growth, herbivore density, and molecular signals. We extend the stochastic Equation 2 to include MEMS input, denoted by a new variable $M(t)$, representing MEMS sensor feedback,

$$dy_1 = \left(a_1 y_1 \left(1 - \frac{y_1}{K} \right) - a_2 y_1 y_2 + M_1(t) \right) dt + \sigma_1 y_1 dW_1,$$

$$dy_2 = \left(a_3 y_1 y_2 - a_4 y_2 - a_5 y_3 y_2 + M_2(t) \right) dt + \sigma_2 y_2 dW_2,$$

$$dy_3 = \left(a_6 y_1 y_2 - a_7 y_3 - a_8 y_3 y_4 + M_3(t) \right) dt + \sigma_3 y_3 dW_3,$$

$$dy_4 = \left(a_9 y_1 y_2 - a_{10} y_4 - a_{11} \cdot \ln_{BIK1} + M_4(t) \right) dt + \sigma_4 y_4 dW_4.$$

For each variable $y_i(t)$, the MEMS feedback terms $M_i(t)$ are defined as follows,

$$M_1(t) = \beta_1 \cdot \text{ControlFunction for } y_1(t),$$

$$M_2(t) = \beta_2 \cdot \text{ControlFunction for } y_2(t),$$

$$M_3(t) = \beta_3 \cdot \text{ControlFunction for } y_3(t),$$

$$M_4(t) = \beta_4 \cdot \text{ControlFunction for } y_4(t).$$

The parameters $\beta_1, \beta_2, \beta_3$, and β_4 represent scaling factors that control the strength of the feedback applied by the MEMS sensors for each variable. Figure 9 presents the results incorporating MEMS feedback into the system. In 5, the plant biomass y_1 shows a significant oscillatory behavior in the stochastic case, with a generally increasing trend after an initial decline. The MEMS feedback helps stabilize biomass growth, resulting in quicker recovery compared to the deterministic solution, which shows a smoother, slower increase. In 5, the insect herbivore density y_2 demonstrates a steady decline, with MEMS further suppressing insect growth. The stochastic model exhibits more variability and faster suppression of herbivores, while the deterministic model shows a smoother, more gradual reduction. The PAD4 protein dynamics y_3 in five show an initial peak followed by a decline and eventual stabilization, with MEMS feedback leading to more active and pronounced fluctuations in the stochastic case compared to the smoother deterministic curve. In 5, the BIK1 protein y_4 follows a similar trend to the other variables, with MEMS inducing stronger oscillations in the stochastic model, particularly towards the later stages. The deterministic solution, on the other hand, stabilizes more smoothly without such fluctuations. The implementation of MEMS introduces dynamic feedback into the system, leading to faster recovery, better control of herbivore density, and more pronounced fluctuations in protein levels. This highlights MEMS' effectiveness in dynamically regulating plant-insect interactions and molecular signals.

6 Discussion

Our comparison of noise conditions revealed distinct behaviors in plant biomass and insect herbivore density. Deterministic conditions produced smooth trajectories, while Gaussian noise caused moderate fluctuations, and Lévy noise led to extreme variations, effectively capturing sudden changes in biological systems. High PAD4 levels and low BIK1 levels were associated with increased plant biomass and reduced herbivore density, indicating effective plant defense mechanisms. Moderate noise intensity ($\sigma = 0.05$) sustained PAD4 activity, providing a balance between system stability and variability.

Lévy noise had a significant impact on PAD4 and herbivore density, with moderate noise intensities sustaining PAD4 activity for optimal plant defense. Queir plots revealed non-linear regulatory interactions between PAD4 and BIK1, emphasizing the role of stochastic dynamics in biological networks. The

results suggest that moderate noise intensity is optimal for maintaining system function, with MEMS-based feedback showing potential for adaptability by continuously adjusting key parameters.

7 Conclusion

This study provides a comprehensive analysis of the dynamic interactions between plants and insect herbivores, focusing on the molecular interplay between PAD4 and BIK1 proteins, under various control strategies and noise conditions. The inclusion of different noise types in the model reveals significant insights into the system's variability. Gaussian noise introduces moderate fluctuations around the deterministic trends, while Lévy noise induces more extreme fluctuations and variability, capturing the sudden changes and extreme events often observed in biological systems. This emphasizes the necessity of incorporating stochastic elements, particularly Lévy noise, to accurately model and understand the complex dynamics of plant-insect interactions. By incorporating MEMS-driven adaptive feedback, we demonstrated how real-time sensor-based interventions can enhance the system's stability and effectiveness, particularly in maintaining plant health and controlling insect populations under varying noise conditions. Our study advances our understanding of plant-insect interactions and offers valuable insights into the role of noise and control strategies in modulating system behavior. These findings have implications for the development of effective management strategies in agricultural and ecological contexts, paving the way for more robust and sustainable approaches to mitigating the impacts of insect herbivory on plant health and productivity.

Data availability statement

The original contributions presented in the study are included in the article/[supplementary material](#), further inquiries can be directed to the corresponding authors.

Author contributions

QA: Conceptualization, Methodology, Software, Writing—original draft, Writing—review and editing. XQ: Funding acquisition, Project administration, Resources, Visualization, Writing—review and editing. NA: Conceptualization, Data curation, Investigation, Methodology, Writing—review and editing. ZK: Funding acquisition, Project administration, Resources, Supervision, Validation, Visualization, Writing—review and editing.

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Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Supplementary material

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fphy.2024.1500423/full#supplementary-material>

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