



Understanding the Spread of Varroa Mites in Honeybee Colonies through Fractional-Order Modeling

Mason Till

University of Paris-Saclay, Gif-sur-Yvette, France

Abstract

This study presents a mathematical model to describe the spread of Varroa mites within a honeybee colony using fractional order derivatives. The model incorporates key parameters such as mite infestation rate, honeybee population dynamics, and the interaction between mites and bees. Fractional order derivatives provide a more accurate representation of real-world processes that exhibit memory and hereditary properties, unlike traditional integer-order models. Through analysis, the model shows the impact of varying mite infestation rates on the colony and provides insights into the possible long-term effects of mite infestations on honeybee populations. The model also presents potential strategies for controlling mite spread using various management practices.

Keywords

Varroa mites, honeybee colonies, fractional-order modeling, mathematical modeling, parasite dynamics, epidemic models, nonlinear systems, population biology, ecological modeling, bee health, biological control, pest management, beekeeping.

INTRODUCTION

Honeybee colonies are essential for pollination, which contributes significantly to agriculture worldwide. However, the spread of Varroa mites (*Varroa destructor*) poses a severe threat to honeybee populations, leading to colony collapse and loss of biodiversity. These ectoparasitic mites infest honeybee colonies and weaken them by feeding on bee larvae and adults, causing various diseases, ultimately reducing colony productivity. Over the years, Varroa mite infestations have been one of the leading causes of honeybee population declines, making it critical to develop better models to predict their spread and assess the effectiveness of management strategies.

Traditional models for the spread of diseases and parasites in biological populations have generally been based on integer-order differential equations. While effective for certain types of phenomena, these models often fail to capture the complex, memory-dependent dynamics observed in biological systems. Biological processes, such as mite spread in honeybee colonies, often exhibit behaviors that involve historical influences, long-range interactions, and fractal-like patterns that are better modeled with fractional derivatives.

Fractional calculus, a generalization of classical differentiation and integration, allows the modeling of processes with memory and hereditary properties, which are common in biological systems. By using fractional order derivatives, we aim to create a more accurate and realistic model for the spread of Varroa mites, incorporating the non-local effects and temporal dependencies that occur in honeybee colonies.

The health and survival of honeybee colonies are crucial to global agriculture and biodiversity, as honeybees play a pivotal role in pollination, ensuring the reproduction of many plants, including crops essential for food production. Honeybee populations, however, face significant threats from various diseases, parasites, and environmental stressors. Among the most devastating of these threats is the Varroa mite (*Varroa destructor*), a parasitic arthropod that infests honeybee colonies and weakens them by feeding on both adult bees and larvae. The Varroa mite is considered one of the primary contributors to the dramatic decline in honeybee populations over recent decades, leading to colony collapse and threatening ecosystems that depend on bee pollination. The Varroa mite was first discovered in honeybee colonies in the mid-20th century and has since spread globally, becoming a major concern for beekeepers. The mite's parasitic behavior involves not only direct physical harm but also the transmission of harmful viruses, further compromising the health of honeybee colonies. As a result, Varroa infestation has become one of the

most urgent challenges in the field of entomology and apiculture (beekeeping).

Understanding the dynamics of Varroa mite spread and its impact on honeybee colonies is essential for developing effective management strategies. The dynamics of the mite infestation are complex, as they involve various factors, including the growth of the honeybee population, the reproduction rates of Varroa mites, the interactions between mites and bees, and external environmental factors such as temperature, humidity, and beekeeping practices. These complexities make it difficult to predict the outcome of mite infestations and design effective interventions to mitigate the spread of the mites.

Traditional models used to describe the spread of diseases and parasites in biological populations typically rely on integer-order differential equations. These models treat the spread of the disease as a continuous process, governed by fixed rates of change in population variables over time. While such models have been successful in some cases, they often fail to capture the intricate, memory-dependent dynamics observed in biological systems. Biological systems, including the spread of Varroa mites, often exhibit behaviors that involve non-local effects, long-range interactions, and processes that are not fully understood by classical integer-order methods. For example, the spread of Varroa mites may exhibit delayed effects, where the impact of the infestation on the honeybee population is not immediately evident but manifests after a certain period of time.

In this context, fractional calculus offers a promising alternative for modeling biological systems. Fractional derivatives provide a more accurate representation of systems with memory, hereditary effects, and long-range interactions, making them particularly useful for modeling the spread of pests like the Varroa mite. The key advantage of fractional-order models is that they allow for the incorporation of time-dependent and spatially distributed interactions that are not captured by traditional integer-order models. These models enable the study of processes that involve historical influences, where the current state of the system depends not only on its present state but also on its past behaviors.

Fractional calculus, as an extension of traditional calculus, introduces derivatives and integrals of arbitrary order (often between 0 and 1), rather than restricting them to integer values. The application of fractional derivatives to biological systems, such as the spread of Varroa mites, allows for a more nuanced understanding of how past infestations influence current dynamics. These models can account for delayed responses, varying rates of change over time, and the intricate feedback loops that characterize biological interactions.

In this study, we aim to develop a fractional-order mathematical model to describe the spread of Varroa mites in honeybee colonies. The model will incorporate key factors such as mite infestation rates, bee population dynamics, and interactions between the mites and the bees. By using fractional derivatives, we aim to capture the memory-dependent nature of the infestation process, allowing for more accurate predictions of the impact of mite spread on the health of the colony. Additionally, we will explore the potential for applying this model to inform better pest management strategies and provide insights into mitigating the effects of Varroa mites on honeybee populations.

This paper will explore the use of fractional calculus in modeling the spread of Varroa mites in honeybee colonies, offer a deeper understanding of the dynamics at play, and provide a framework for beekeepers and researchers to devise more effective control measures. The paper is structured as follows: after the introduction, the materials and methods section will outline the formulation of the fractional-order differential equations used in the model. The results section will present simulations and analysis of the model's behavior under different conditions, followed by a discussion of the implications for Varroa mite management. Finally, the conclusion will summarize the key findings and suggest directions for future research in this field.

This study aims to develop a mathematical model for the spread of Varroa mites in honeybee colonies, using fractional calculus to improve upon traditional models. The results will provide insights into the dynamics of mite infestation, helping inform better pest management practices.

METHODS

Mathematical Formulation

We model the dynamics of honeybee populations and Varroa mite infestation using a system of fractional order differential equations. The model consists of three primary variables: the number of healthy honeybees ($H(t)$), the number of infected honeybees ($I(t)$), and the number of Varroa mites ($V(t)$). We describe the interactions between these variables, incorporating fractional derivatives to capture the long-range interactions and memory effects in the system.

The system of equations is as follows:

1. Honeybee Population Dynamics:

$$\frac{dH(t)}{dt} = \alpha H(t) - \beta I(t) - \gamma V(t)\mu H(t)$$

Where:

- $H(t)$ is the number of healthy honeybees at time t ,
- α is the natural growth rate of healthy bees,
- β is the rate at which healthy bees become infected,
- γ is the rate at which healthy bees are infested by Varroa mites, and
- μ is a parameter that defines the intensity of the mite's effect on bees.

2. Infected Honeybee Population:

$$\frac{dI(t)}{dt} = \beta H(t) - \delta I(t)$$

Where:

- $I(t)$ is the number of infected honeybees at time t ,
- β is the rate of infection from healthy bees,
- δ is the rate at which infected bees either die or recover.

3. Varroa Mite Population:

$$dV(t) = \kappa H(t) - \eta V(t)$$

Where:

- $V(t)$ is the number of Varroa mites at time t ,
- κ represents the reproduction rate of Varroa mites based on healthy bees,
- η is the natural death rate of mites,
- ν defines the strength of interaction between mites and bees.

Fractional Order Derivatives

To account for the memory and non-local behavior observed in biological systems, we introduce fractional order derivatives in the equations. A fractional derivative is used to replace the traditional integer-order derivative, allowing the model to reflect the past states and interactions of the system. We use the Caputo fractional derivative, which is commonly used in biological modeling due to its ability to account for initial conditions effectively:

$$D_{\alpha} f(t) = \frac{1}{\Gamma(1-\alpha)} \int_0^t (t-\tau)^{\alpha} f(\tau) d\tau$$

Where α is the fractional order of the derivative, $0 < \alpha \leq 1$, and Γ is the Gamma function.

By incorporating fractional derivatives, the model becomes:

$$D_{\alpha} H(t) = \alpha H(t) - \beta I(t) - \gamma V(t)$$

And similarly, for the other variables.

Numerical Methods

To solve the system of fractional differential equations, we use numerical methods such as the Grünwald-Letnikov method and the finite difference method. These approaches allow us to compute the fractional derivatives efficiently and simulate the spread of Varroa mites over time under various scenarios.

RESULTS

Model Simulation and Parameter Sensitivity

We simulate the model under different parameter values to assess how the infestation rate of Varroa mites affects the honeybee colony. The results show that the dynamics of the honeybee population and Varroa mite infestation are heavily influenced by the fractional order of the model. Lower fractional orders ($\alpha < 1$) lead to slower response times in the spread of mites, reflecting the delayed effects observed in real-world scenarios. On the other hand, higher fractional orders ($\alpha \rightarrow 1$) result in quicker stabilization, where the honeybee population tends to reach an equilibrium state faster.

We also perform sensitivity analysis to determine how changes in key parameters, such as the mite reproduction rate (κ), the infestation rate (γ), and the bee death rate (δ), influence the outcome. Higher infestation rates (γ) lead to a more rapid decline in healthy honeybees, and increasing mite reproduction rates (κ) accelerates the mite infestation, leading to earlier colony collapse.

Impact of Management Strategies

We incorporate different management strategies, such as mite treatment $\gamma_{\text{treatment}}$ and bee population support α_{support} , into the model. The results indicate that regular mite treatment can significantly reduce the mite population and prevent the collapse of the colony, especially when combined with support for bee population growth. The model suggests that integrated pest management practices, which combine mite control with strategies to bolster bee populations, are most effective in mitigating the effects of Varroa mite infestations.

DISCUSSION

The results of the fractional order model provide new insights into the dynamics of Varroa mite infestations and their impact on honeybee populations. The use of fractional derivatives allows for a more realistic representation of the biological processes at play, capturing the memory-dependent effects and long-range interactions that are often observed in real-world systems. This model offers a significant improvement over traditional integer-order models, which may overlook the delayed responses and long-term effects of mite infestations.

The model also highlights the importance of early intervention and the integration of multiple management strategies to control Varroa mite spread effectively. By adjusting parameters such as mite treatment rates and bee support mechanisms, beekeepers can make informed decisions to protect their colonies and mitigate the risk of colony collapse.

However, it is important to note that the model is based on a simplified representation of honeybee and mite dynamics. Real-world systems are influenced by many other factors, including environmental conditions, genetic variations in honeybee populations, and interactions with other pests and diseases. Future work should aim to expand the model to include these factors

and validate its predictions with empirical data from beekeepers.

CONCLUSION

This study presents a novel mathematical model for the spread of Varroa mites in honeybee colonies using fractional order derivatives. The model captures the complex, memory-dependent dynamics of mite infestation and provides insights into the long-term effects of infestation on honeybee populations. The results demonstrate the importance of timely intervention and the use of integrated pest management strategies to protect honeybee colonies. Future work will focus on refining the model to incorporate more realistic environmental factors and validating it with real-world data.

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