

REVIEW

Biotic and abiotic stresses on honeybee health

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Abstract

Honeybees are the most critical pollinators providing key ecosystem services that underpin crop production and sustainable agriculture. Amidst a backdrop of rapid global change, this eusocial insect encounters a succession of stressors during nesting, foraging, and pollination. Ectoparasitic mites, together with vectored viruses, have been recognized as central biotic threats to honeybee health, while the spread of invasive giant hornets and small hive beetles also increasingly threatens colonies worldwide. Cocktails of agrochemicals, including acaricides used for mite treatment, and other pollutants of the environment have been widely documented to affect bee health in various ways. Additionally, expanding urbanization, climate change, and agricultural intensification often result in the destruction or fragmentation of flower-rich bee habitats. The anthropogenic pressures exerted by beekeeping management practices affect the natural selection and evolution of honeybees, and colony translocations facilitate alien species invasion and disease transmission. In this review, the multiple biotic and abiotic threats and their interactions that potentially undermine bee colony health are discussed, while taking into consideration the sensitivity, large foraging area, dense network among related nestmates, and social behaviors of honeybees.

Key words: abiotic stressor, *Apis*, biotic stressor, colony losses, interaction effect, pollinator decline

INTRODUCTION

Pollinators provide an important ecosystem service by playing a key functional role in sustaining both wild plant communities and agricultural productivity (Potts *et al.* 2016). Animal-mediated pollination services benefit over three-quarters of crop plants (Porto *et al.* 2020), and the immense value of insect pollination in improving crop quantity and quality has been widely recognized (Fijen *et al.* 2018; Khalifa *et al.* 2021). Insects, particularly honeybees, are the primary pollinators of both crops and wild plants. However, concerns have been raised that invertebrate pollinator populations, both in the wild

and under management, are declining worldwide, particularly in the Northern Hemisphere (Neumann & Carreck 2010; Pirk *et al.* 2016; Ulrich *et al.* 2020; van der Sluijs 2020). Although this argument has been sometimes challenged (e.g. Ghazoul 2005), the dependency of crops on pollinators is undeniably increasing (Aizen *et al.* 2019; Garibaldi *et al.* 2020), and alternatives such as robot bees cannot yet effectively counter pollination deficiency (Potts *et al.* 2018; Gleadow *et al.* 2019). In many parts of the world, reliance on managed pollinators, including the western and eastern honeybees—*Apis mellifera* and *Apis cerana*, respectively, is prevailing owing to the value of their products and their crop pollination efficiency both spatially and in a timely manner. Honeybees live in colonies and visit various flowering plants by foraging, during which they face a variety of biotic and abiotic threats and frequently interact with related nestmates and intercolonial or interspecies individuals (Leong

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et al. 2016; Quigley *et al.* 2019; Pusceddu *et al.* 2021; Siviter *et al.* 2021). Consequently, colony loss resulting from the aforementioned challenges has emerged as a crucial issue in global apiculture (Steinhauer *et al.* 2018; Hristov *et al.* 2020). In this review, we aimed to comprehensively summarize the multiple biotic and abiotic threats and their interactions that potentially undermine bee colony health to offer insights regarding the studies of honeybee health and to provide guidance to the apiculturists and policymakers.

BIOTIC STRESSORS

Ectoparasitic mites

Two ectoparasitic mite genera, *Varroa* and *Tropilaelaps* with a global and mostly Asian distribution, respectively, inflict insurmountable problems on *A. mellifera* populations (Nazzi & Le Conte 2016; de Guzman *et al.* 2017). Although four species of each *Varroa* and *Tropilaelaps* have been identified, only *Varroa destructor* and *Tropilaelaps mercedesae* are threatening colony health, to an extent that could lead to colony collapse if not for human intervention (Chantawannakul *et al.* 2018; Traynor *et al.* 2020). This is evident from the fact that most wild and feral populations of *A. mellifera* in the Northern Hemisphere have been eradicated due to the lack of treatment (Kraus & Page 1995; Oldroyd 2007; Strauss *et al.* 2016; Pirk *et al.* 2017).

Varroa destructor feeds exclusively on the host's fat body, consuming nearly a microliter daily (Ramsey *et al.* 2018, 2019; Posada-Florez *et al.* 2019) and significantly interfering with honeybee physiology (Amdam *et al.* 2003; Arrese & Soulages 2010). The life cycle of *V. destructor* consists of two stages, the phoretic dispersal phase, during which it parasitizes adult bees, and the reproductive phase, closely synchronized with the host's development, during which it infests immature individuals (Traynor *et al.* 2020).

Most biological features of *Tropilaelaps* are similar to those of *Varroa* excluding the former's smaller body size, rapid locomotion, faster reproduction rate, and ability to mate outside brood cells (de Guzman *et al.* 2017; Chantawannakul *et al.* 2018; Gao *et al.* 2021). Although *Tropilaelaps* mites are more prevalent than *Varroa* mites in some Asian *A. mellifera* colonies (Buawangpong *et al.* 2015), studies on this parasitic mite are still in their initial stages, and more investigations are needed.

Several behaviors, such as worker auto-grooming, allo-grooming, and hygienic behavior (Bak & Wilde 2015; Locke 2016; Leclercq *et al.* 2017; Nganso *et al.* 2017; Pusceddu *et al.* 2021), have been observed in naturally surviving *A. mellifera* colonies, whereas reduced mite-reproductive success seems to be a primary determinant of mite-resistance (Oddie *et al.* 2017; Nganso *et al.* 2018). Notably, cell recapping was recently evidenced to be associated with lower mite reproductive success (Lin *et al.* 2018; Grindrod & Martin 2021; Oddie *et al.* 2021). Indeed, controlling the reproduction of parasites is key to maintaining the balance of the host-parasite relationship in favor of the host. Therefore, investigating the population dynamics of ectoparasitic mites and their host-finding and reproductive behaviors is vital for controlling these mites in honeybee colony management. Once the directional mechanism of mites is determined, biological approaches, such as pheromones for trapping, repelling, or interfering with their mating behavior, can be used to limit the population growth of these ectoparasites.

A lack of coevolution between ectoparasitic mites and *A. mellifera*, which was first infested a few decades ago (De Jong *et al.* 1982), is considered a key factor in the susceptibility of the honeybee host. Considering the poor efficacy and negative side effects of medication-based approaches (Dietemann *et al.* 2012), breeding efforts are aimed at selecting honeybee stocks that can resist these serious biotic threats worldwide (Büchler *et al.* 2010). The immune mechanism responsible for the defense of bee colonies and for individuals to cope with external threats is characterized by plasticity (Mondet *et al.* 2015), and thus, artificially selecting honeybee stocks (Hopkins *et al.* 2012) for resistance to primary biotic threats is an important form of human intervention to accelerate colony evolution (Lin *et al.* 2023) and achieve resistance. However, relevant studies have made limited progress in recent years due to, in our opinion, knowledge gaps in the relationship between mites and their original hosts. Recent evidence supports that immature *A. cerana* individuals are more susceptible (Page *et al.* 2016; Lin *et al.* 2018, 2022) and its adult workers are significantly more efficient in the rapid detection and removal of dead brood (Lin *et al.* 2016), which is also observed in the mite-resistant *A. mellifera* stocks (Villa *et al.* 2009), limiting the reproduction of parasitic mites (Page *et al.* 2016; Lin *et al.* 2018, 2021a; Wang *et al.* 2020). These characteristics can be informative for *A. mellifera* breeding programs.

Microorganisms

Pathogenic microorganisms play a crucial role in ecosystems (Preston *et al.* 2016; Piot *et al.* 2022). Host pathogen infections can be divided into covert and overt ones, determined by factors such as the coevolution time between the pathogen and host, physiological function of the host, antiviral response, transmission route, and environmental factors (Rigaud *et al.* 2010; Britton & Jane White 2021). In covert infections, the proliferation of pathogens is limited, reducing the probability of the horizontal transmission of pathogens between hosts. However, the long-term survival of infected hosts increases the likelihood of vertical transmission from queens and drones to offspring (Chen *et al.* 2006b). In contrast, overt infections are typically characterized by the rapid replication of the pathogen within an individual, leading to increased virulence and opportunities for horizontal transmission via trophallaxis and shared food resources (Chen *et al.* 2006a).

Owing to the rapid advances in and increasing accessibility of high-throughput sequencing technologies, >60 honeybee-infecting viruses have been identified (Beaurepaire *et al.* 2020), with the most detrimental being mainly single-stranded positive-sense RNA viruses from the families of *Dicistroviridae* and *Iflaviridae*. Viruses from the *Dicistroviridae* family (dicistroviruses) include acute bee paralysis virus, black queen cell virus, Israeli acute paralysis virus, and Kashmir bee virus, whereas *Iflaviridae* viruses (iflavirus) include deformed wing virus, slow bee paralysis virus, and sacbrood virus, as well as the taxonomically unsystematic chronic bee paralysis virus (Grozinger & Flenniken 2019; Procházková *et al.* 2020; Ullah *et al.* 2021). Sacbrood virus and deformed wing virus, in particular, have attracted the most attention and have been the subject of investigation in *A. cerana* and *A. mellifera* virology, respectively (Gong *et al.* 2016; Beaurepaire *et al.* 2020; Wei *et al.* 2022). The structures of most of these common viruses have been deciphered (Kalynych *et al.* 2017; Škubník *et al.* 2017, 2021; Spurný *et al.* 2017; Procházková *et al.* 2018; Acosta-Reyes *et al.* 2019), aiding the development of antiviral compounds. Both dicistroviruses and iflavirus are enveloped in capsids with icosahedral symmetries, while iflavirus genome release is facilitated by a large opening of the viral capsid and triggered by an acidic pH, which is typical in endosomes, indicating that the viruses may enter cells by endocytosis (Procházková *et al.* 2020; Škubník *et al.* 2021). However, our understanding of honeybee virology at the molecular level is limited because of a lack of stable honeybee tissue cell lines (Guo *et al.* 2020).

Similar to other organisms, bee viruses affect various aspects of honeybee biology and cause a variety of well-recognized effects, such as body malformation, stunted development, tremor paralysis, decreased immunity, diminished reproductive capability, impaired cognition and homing ability, navigation system loss, and shortened life span (McMenamin & Flenniken 2018; reviewed by Ullah *et al.* 2021). The spread and virulence of viruses can be largely facilitated by ectoparasitic mites (discussed in the section of Interactive and cumulative effects of multiple stressors).

In addition to viruses, bacterial (e.g. *Paenibacillus larvae* and *Melissococcus plutonius*) and fungal (e.g. *Nosema ceranae*) pathogens threaten honeybee health. *Paenibacillus larvae* and *M. plutonius*, the causative agents of American foulbrood and European foulbrood, mainly infect the larvae of *A. mellifera* and *A. cerana*, respectively, consequently leading to the collapse of a colony if left untreated (Alonso-Salces *et al.* 2017; de León-Door *et al.* 2020; Devi *et al.* 2021). Although the absence of antibodies in insects leads to a deficiency in transgenerational immunity, the survival rate of the offspring of orally vaccinated queens can be largely increased after *P. larvae* challenge (Dickel *et al.* 2022). *Nosema ceranae* is an obligate intracellular eukaryotic parasite that affects adult bees and has been intensively studied over the past dozen years since its first detection. It was extensively detected coincidentally in collapsing colonies after infection similar to the outbreak of Colony Collapse Disorder scenario in the United States (Chen *et al.* 2008; Higes *et al.* 2008; Martín-Hernández *et al.* 2018). *Nosema ceranae* invades the midgut and affects different castes of several bee species (Goulson *et al.* 2015) in terms of physiological processes, behavioral performance, metabolism, immune response, and so on (Paris *et al.* 2018; reviewed by Martín-Hernández *et al.* 2018).

Large biotic enemies

Honeybees face numerous biotic enemies during nest-living and foraging activities, such as the predatory hornet, wax moth, mantis, Batrachia, and small hive beetle. The giant hornet *Vespa*, causing honeybee foragers homing failure and colony foraging paralysis, is native to Asia and invaded Europe and America during this century (Requier *et al.* 2019; Zhu *et al.* 2020; Werenkraut *et al.* 2022). This predator hunts arthropods, including *A. cerana* and *A. mellifera*, for food and decreases the survival probability of colonies of the latter due to a lack of effective defenses (Tan *et al.* 2007; Requier *et al.* 2019).

The small hive beetle (*Aethina tumida*), feeding on both fruits and bee nest products, originated in sub-Saharan Africa and is now distributed almost globally (Evans *et al.* 2018), which may partially be facilitated by the international beeswax trade (Idrissou *et al.* 2019). At present, our limited knowledge of its biology, diagnosis, control, and prevention restricts the development of efficient treatments (reviewed by Neumann *et al.* 2016), which is also the case for other honeybee pests and diseases. Therefore, maintaining a healthy colony via good beekeeping management is commonly the ideal approach for disease prevention.

ABIOTIC STRESSORS

Agrochemicals

In modern agricultural systems, pollinators are readily exposed to agrochemicals during periods that they heavily rely on blooming crops. Honeybees can be utilized as bioindicators of the effects of agrochemicals and land-use on pollinators in view of their sensitivity and ability to provide high-resolution information regarding the presence of environmentally persistent agrochemicals and to reflect changes in agricultural landscape quality at spatial and temporal scales (Quigley *et al.* 2019). Mounting evidence suggests that agrochemicals, including pesticides, fungicides, herbicides, and acaricides, are one of the most direct and severe threats to honeybee physiology and colony health, and some of them, particularly neonicotinoids, have been directly implicated in colony collapse (Goulson *et al.* 2015; Sánchez-Bayo *et al.* 2016; Siviter *et al.* 2021; Wang *et al.* 2021a). Numerous types of extremely toxic agrochemicals that directly cause acute toxicity and lethality to pollinators are banned or restricted in agricultural practices in several countries. However, the sublethal negative effects of medium- or low-toxicity agrochemicals, partially due to their prophylactic use in the surroundings, can still disrupt the physiology and behavior of all caste bees in the hive.

The reproductive toxicity of agrochemicals to queens and drones has always been in focus as it dramatically affects colony growth and sustainability. Even in cases of high mating frequencies, neonicotinoid-exposed queens may have compromised ovaries and spermathecal-stored sperm quality and quantity, adversely affecting reproduction (Rangel & Tarpy 2015; Williams *et al.* 2015). Physiological disorders in the queens, such as decreased attractiveness of pheromones to workers (Walsh *et al.* 2020) and dysfunctional gene expression related to an-

tioxidant, immunity, and development, lead to reduced control of the colony (Chaimanee *et al.* 2016; Costa *et al.* 2022). Agrochemicals can significantly affect the quality and viability of drones and their sperm (Straub *et al.* 2016). Deltamethrin, acetamiprid, fumagillin, thiamethoxam, clothianidin, and fipronil exposure reduces drone survival and sperm viability and concentration and increases the percentage of sperm with defective membrane integrity while increasing sperm metabolism, which may be associated with drone infertility (Kairo *et al.* 2016; Fisher & Rangel 2018; Ben Abdelkader *et al.* 2021; Straub *et al.* 2021). These stresses on queens and drones undermine sustainability, directly resulting in the loss of workers and causing colony collapse.

Worker bees constitute the largest part of the population and undertake almost all labor in hives. Compared with queens and drones, worker bees are more likely to be exposed to agrochemicals because of their foraging activities. The transfer of agrochemical residues in the hive compartments to the honeybee brood, augmented in both variety and dose, is inevitable (Morales *et al.* 2020). In addition to acute toxicity and death (Raymann *et al.* 2018; Tihelka 2018), sublethal effects of agrochemicals on brood development, including delayed larval development and reduced adult hatching weight, have been observed (Wu *et al.* 2011; Odemer *et al.* 2020), especially under high hive temperatures where the sublethal effects are exacerbated (Medrzycki *et al.* 2010; Wu *et al.* 2011). The dysfunctional development of the neural tissues caused by neurotoxic substances has been extensively investigated in honeybees (Desneux *et al.* 2007). Neonicotinoids have been shown to impair learning and memory in honeybees, directly affecting the ability of workers to navigate and forage (Tan *et al.* 2017; Muth & Leonard 2019; Muth *et al.* 2019). Consequently, workers appear disoriented, have a degenerated sucrose responsiveness (Eiri & Nieh 2012; D'Émara *et al.* 2016), consume less food (Wang *et al.* 2022), and require more time to discern food. These physiological changes may explain why workers exhibit decreased waggle dancing and foraging success after neonicotinoid exposure (Eiri & Nieh 2012; Henry *et al.* 2012; Teeters *et al.* 2012). Additionally, agrochemicals have the potential to disrupt gut microbes (Kakumanu *et al.* 2016; Paris *et al.* 2020; Wang *et al.* 2023) and increase susceptibility to infection by opportunistic pathogens (Motta *et al.* 2018).

Environmental change

A decreased pathogenicity of common pathogens in the wild non-cave *Apis* spp. (Lin *et al.* 2021b) indicates the

potential role of environmental change in the observed decline of pollinators. Amidst the backdrop of global change reflected by climate change, expanding urbanization, and agricultural intensification, pollinator biodiversity and flower-rich habitats are being destroyed or fragmented (Potts *et al.* 2016; Goulson & Nicholls 2022). Climate change affects flora and fauna and their interactions, including flowering phenology, floral rewards, honeybee behavior, and plant pollinator interactions, resulting in novel communities and altered ecosystems (Keeler *et al.* 2021; Vercelli *et al.* 2021). Extreme temperatures and reduced precipitation can end flower blooms prematurely, accelerate the wilting of flowers, and reduce pollen and nectar availability (Flores *et al.* 2019; Sayğı 2020), affecting the symbiosis of honeybees and flowering plants that they pollinate, given that time and pattern of flowering, as well as pollinating activity, are sensitive to temperature (Hegland *et al.* 2009; Forrest 2015). The prolificacy of queens, spermatogenesis of drones, and viability of honeybee broods can also be negatively influenced by high atmospheric temperatures, leading to a reduction in the reproductive function of queens and thermoregulation of worker bees, and thus to colony homeostasis disruption (Sales *et al.* 2018; Cebotari *et al.* 2019; Walsh *et al.* 2019; McAfee *et al.* 2020; Cunningham *et al.* 2022).

Urbanization and land-use intensity can lead to the loss of natural habitats for bees and thus to decreased abundance of floral resources and nesting opportunities, resulting in a reduction in the number, richness, and diversity of bees (Jaffé *et al.* 2019; Millard *et al.* 2021). A decline in brood emergence occurs when honeybees are unable to adapt or find new habitats, affecting colony continuation (McCabe *et al.* 2022). Although urbanization can increase the abundance of floral resources to a certain extent, which would benefit some honeybee populations and partially compensate for its negative effects, this is often limited by urban warming and the loss of agricultural land (Wilson & Jamieson 2019). The expanse of industrial agriculture has intensely modified the landscape across large areas of the world, often reducing the availability and diversity of floral resources (Goulson & Nicholls 2022). As a result of agricultural intensification practices, such as large-scale cultivation, monoculture, and highly disturbed agronomic practices, the biodiversity of pollinating insects in and around fields has been decreased (Marshman *et al.* 2019; Seibold *et al.* 2019; Raven & Wagner 2021). Overall, the impact of environmental change can be determined by studying the accessibility of floral resources and the functional characteristics of honeybee populations.

Environmental pollutants

Honeybees interact with air, soil, water, and utilitarian plants during flight and foraging and are directly exposed to various environmental pollutants, such as microplastics, heavy metals, and particulate matter (PM). Vast quantities of pollutants have been detected in different tissues of pollinators (Feldhaar & Otti 2020; Deng *et al.* 2021; Edo *et al.* 2021). Honeybees are, therefore, generally used as sentinel bioindicators in environmental monitoring plans due to their sensitivity, large foraging area, morphological features, and social behaviors (Bargańska *et al.* 2016; Herrero-Latorre *et al.* 2017; Goretti *et al.* 2020; Gutiérrez *et al.* 2020; Capitani *et al.* 2021). Honeybees could also serve as bioindicators of environmental SARS-CoV-2 prevalence, since coronavirus particles have widely been detected in the PM carried by foraging workers (Cilia *et al.* 2022). In addition, as honeybees distribute environmental pollutants within the colony during food transfer, the risks of environmental pollutants present in bee products to human health also need to be considered (Diaz-Basantes *et al.* 2020; Papa *et al.* 2021; Smith & Weis 2022).

The large-scale use of plastic has contributed to huge quantities of hazardous refuse being produced at a global level and represents one of the most prominent issues of the Anthropocene (Cox *et al.* 2019; Zhang *et al.* 2020). Microplastics are ubiquitous and persistent in almost all environments and pose a potential threat to a variety of plant and animal species (Anbumani & Kakkar 2018; Mammo *et al.* 2020). Exposure to microplastics will commonly not lead to acute mortality of honeybees in the short term; rather, it decreases the gut microbiome diversity, alters the microbiome structure, and changes the expression of genes related to detoxification and immunity (Wang *et al.* 2021b; Balzani *et al.* 2022; Buteler *et al.* 2022).

Atmospheric PM, primarily emitted by vehicular transport, combustion engines, coal mining, and agricultural residues, have been demonstrated to affect the navigation of honeybee workers, increasing the foraging duration, and affecting bee survival, flower visitation, heart rate, hemocyte levels, and gene expression (Thimmegowda *et al.* 2020; Cho *et al.* 2021). Coupled with heavy metals, such as lead, copper, and manganese, the exposures exert potential neurotoxic effects, including damage to learning and long-term memory formation through impaired protein functioning, gut lining damage and microbiome alterations, immune system activation, and aberrant DNA methylation, eventually causing lethal or sublethal effects on honeybees (Burden 2016; Costa *et al.* 2019;

Al Naggar *et al.* 2020). Furthermore, noise, turbulence and dust in road verges have been evidenced to impact honeybee forager activities as well (Dargas *et al.* 2016; Phillips *et al.* 2021), although the benefits of road verges to pollinators may outweigh the costs (Phillips *et al.* 2020).

Human management

Lack of natural selection

Although beekeepers play a crucial role in controlling colony diseases, such as regular queen renewal to promote honeybee health (Simeunovic *et al.* 2014), the role of apiculture in limiting natural selection and evolution negatively contributes to colony health (Neumann & Blacquière 2017; Oddie *et al.* 2017). To maintain a productive colony, modern beekeepers perform regular treatments to ensure its survival, which is contrary to the law of natural selection (Le Conte *et al.* 2020). Drug treatments may interfere with the natural population dynamics of colonial endosymbionts (Neumann & Blacquière 2017). In addition, the intensification of colony density by commercial beekeeping increases the probability of drifting foragers, thus facilitating the pathophoresis of infectious diseases (Seeley & Smith 2015).

Interspecific competition and disease transmission after colony translocations

The anthropogenic introduction of domestic species to regions beyond their natural habitat range can lead to competition with endemic creatures and may result in biological invasions affecting living organisms (Young *et al.* 2017). Although there is little evidence that introduced managed honeybees have a major impact on the survival, fecundity, and population density of other pollinators, they most likely have negative effects on other honeybee species (Moritz *et al.* 2005). The high number of introduced *A. mellifera* colonies in Asia could lead to direct competition with endemic cavity-nesting honeybee species (such as *A. cerana*, *Apis koschevnikovi*, *Apis nigrocincta*, and *Apis nuluensis*) for nest sites and with all managed and wild honeybees for floral resources (IPBES 2016; Geslin *et al.* 2017; Mallinger *et al.* 2017; Vanbergen *et al.* 2018). *Apis mellifera* may also interfere with mating of virgin *A. cerana* queens and steal stored honey from native honeybees (Theisen-Jones & Bienefeld 2016).

It is widely recognized that the introduction of alien species can lead to the spread of infectious diseases (Crowl *et al.* 2008; Hulme 2014). Managed honeybee

translocations across regions, countries, and continents facilitated by human migration and trade have contributed to ongoing disease transmission as a consequence of colony introduction and communication (Beaurepaire *et al.* 2020). Meanwhile, alien species in a new habitat are more susceptible to diseases than native pollinators, which usually have a survival advantage, as demonstrated by the greater susceptibility of *A. mellifera*, introduced in Southeast Asia, to the invasive ectoparasitic *Varroa* and *Tropilaelaps* mites compared to that of the native *A. cerana* (Nazzi & Le Conte 2016; de Guzman *et al.* 2017; Traynor *et al.* 2020). In this regard, the invasion of alien mites has constrained the establishment of feral *A. mellifera* populations in Asia so far. In other words, ectoparasites can protect wild and managed Asian honeybees, including their original hosts against invasion.

Malnutrition

Adequate nutrition is fundamental to the health of organisms at all life stages (Wickramasinghe *et al.* 2020). Nectar and pollen, the primary nutritional resources collected by honeybees, are rich in carbohydrates, proteins, lipids, and micronutrients (Vaudou *et al.* 2015; Tsuruda *et al.* 2021). Apicultural practices govern colony nutrition through the choice of foraging areas for worker bees (Neumann & Blacquière 2017). However, the increased colony density in modern apiculture leads to high demand for foraging resources. Beekeepers exploiting honeybee colonies for hive products and providing workers with syrup and soybean flour in exchange can compromise colony health and affect its resilience to diseases (Mao *et al.* 2013; Wheeler & Robinson 2014). Evidence emerges that a modest amount of probiotics appears beneficial to the fitness of honeybees (Evans & Lopez 2004; Brown *et al.* 2022), although few studies have investigated the issue despite the key role of the gut microbiota in bee health (Kwong & Moran 2016). Unfortunately, because of the nutritional knowledge gap, a standard balanced diet is currently lacking in commercial beekeeping, highlighting the need for further studies to correct this situation (Paray *et al.* 2021; reviewed by Tsuruda *et al.* 2021).

INTERACTIVE AND CUMULATIVE EFFECTS OF MULTIPLE STRESSORS

Generally, biotic and abiotic stressors do not act independently, and quite often, additive or synergistic effects have been demonstrated to impact colony health (O’Neal *et al.* 2018; Siviter *et al.* 2021; Goulson & Nicholls 2022). Typically, there is a finely balanced relationship

between microbial pathogens and their host (Varela *et al.* 2009; Deroost *et al.* 2015). Multiple microbial pathogens can interact with each other to achieve mutual benefit and reciprocity (Gajda *et al.* 2021). However, when the natural dynamic balance is disturbed, the pathogens can inflict devastating effects on their hosts. For example, pathogen virulence can increase with the introduction of a second host when a multi-host pathogen, such as a deformed wing virus vectored by *V. destructor*, spreads among species (Woolhouse *et al.* 2001; Martin *et al.* 2012). *Varroa destructor* can serve as a harbor for virus replication and transmission as a result of bias proliferating highly virulent strains and weakening host immunity (Di Prisco *et al.* 2011; Martin *et al.* 2012; Lin *et al.* 2022). Furthermore, virus-induced immunosuppression in hosts can favor the feeding and reproductive behavior of ectoparasitic mites (Di Prisco *et al.* 2016). On the other hand, the fitness of mites may also be influenced by the vectored viruses, a subject yet to be further investigated (e.g. Wang *et al.* 2019; Ryabov *et al.* 2022).

The decline in honeybee populations has been reported to be associated with nutritional shortage caused by land-use intensification (Vaudo *et al.* 2015). A well-sustained nutritional state not only enhances the survival of pathogen-infected hosts, but also significantly shapes the microsporidian–virus interface (Zheng *et al.* 2014; Tritschler *et al.* 2017). On the contrary, nutritional stress adversely affects honeybee immunity and renders bees more susceptible to microbial pathogens, while vice versa some pathogens have detrimental effects on the nutritional resilience of their hosts (Dolezal & Toth 2018; Grozinger & Flenniken 2019; Castelli *et al.* 2020). Additionally, nutritional stress combined with pesticides can reduce the food consumption of honeybees and their flight success, leading to increased mortality (Tong *et al.* 2019). Regarding other abiotic factors, urbanization increases pathogenic pressure on honeybees, and climate change will increase the risk of hornet and non-native bee species invasions (Barbet-Massin *et al.* 2013; Tabor & Koch 2021) and affect the prevalence of infectious diseases, including viruses, American foulbrood, *N. ceranae*, and small hive beetles (Lindgren *et al.* 2012; Martín-Hernández *et al.* 2018; Cornelissen *et al.* 2019; Jamal *et al.* 2021; de Jongh *et al.* 2022; Piot *et al.* 2022). The ingestion and accumulation of microplastics and antibiotics in honeybees can also undermine their antiviral defenses (Deng *et al.* 2021, 2022). In addition to viruses and *Nosema* (Aufauvre *et al.* 2012; Di Prisco *et al.* 2013), pesticide exposure can impair both the immune and detoxification system of honeybees, leading to increased susceptibility to parasites (Straub *et al.* 2019; Annoscia *et al.* 2020).

CONCLUSION

In the context of widespread insect and pollinator decline (Potts *et al.* 2016; McDermott 2021), honeybee health has long been a point of discussion in the apiculture and scientific community (Hassler *et al.* 2021). This critical eusocial pollinator encounters a variety of biotic and abiotic stressors during foraging and nesting. Further research is required to investigate other potential factors that may plausibly affect honeybee physiology, such as electromagnetic radiation (Molina-Montenegro *et al.* 2023). Generally, a combined effect of multiple stressors is likely to be more harmful than a single stressor (Goulson *et al.* 2015; Goulson & Nicholls 2022). It can be argued that the interaction among multiple stressors, varying spatially and temporally, is the key factor underlying the core issue of global honeybee health.

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CONFLICT OF INTEREST

The authors declare no competing financial interests.

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