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MICRO-REVIEW

Microalgae as a promising and sustainable nutrition source for managed honey bees

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Abstract

Managed honey bee colony losses are attributed to a number of interacting stressors, but many lines of evidence point to malnutrition as a primary factor. Commercial beekeepers have become increasingly reliant on artificial pollen substitute diets to nourish colonies during periods of forage scarcity and to bolster colony size before pollination services. These artificial diets may be deficient in essential macronutrients (proteins, lipids, prebiotic fibers), micronutrients (vitamins, minerals), and antioxidants. Therefore, improving the efficacy of pollen substitutes can be considered vital to modern beekeeping. Microalgae are prolific sources of plant-based nutrition with many species exhibiting biochemical profiles that are comparable to natural pollen. This emerging feed source has been employed in a variety of organisms, including limited applications in honey bees. Herein, I introduce the nutritional value and functional properties of microalgae, extrapolating to central aspects of honey bee physiology and health. To conclude, I discuss the potential of microalgae-based feeds to sustainably provision managed colonies on an agricultural scale.

KEYWORDS

antioxidant, *apis mellifera*, microbiota, pollen substitute, prebiotic

1 | INTRODUCTION

Nutritional status is among the most important modifiable determinants of honey bee health. In honey bees, malnutrition interacts with and amplifies the detrimental effects of parasites, pathogens, and pesticide exposure (Dolezal & Toth, 2018; Dolezal et al., 2019; Tosi, Nieh, Sgolastra, Cabbri, & Medrzycki, 2017). Pollen is the colony's main source of proteins, lipids, vitamins and minerals (Brodtschneider & Crailsheim, 2010). As such, beekeepers

bioavailable and provide all essential amino acids (EAAs), which are present in almost half its total protein content (Soni, Sudhakar, & Rana, 2017). *Chlorella* are green microalgae that contain 50–60% protein dry weight and a full complement of EAAs (Kotrbaček, Doubek, & Doucha, 2015). The EAAs required by honey bees are arginine, histidine, isoleucine, leucine, lysine, methionine, phenylalanine, threonine, tryptophan, and valine (De Groot, 1953), all of which must be obtained from pollen. When nurse bees consume pollen, amino acids are incorporated into the fat body storage lipoprotein *vitellogenin*, which acts as a precursor to proteinaceous hypopharyngeal gland secretions (jelly) used to rear brood (Crailsheim, 1990). The microalga *Chlorella sorokiniana* was recently shown to positively impact honey bee colony growth and individual nutritional physiology (fat body protein content, *vitellogenin* messenger RNA levels, and hypopharyngeal gland size), indicating that *Chlorella* is promising as a PS or diet additive (Jehlík et al., 2019). In one study, *Arthrospira* was incorporated into a PS formulation that led to improved colony performance compared to unfed control colonies (Kumar, Mishra, & Agrawal, 2013), however the effects of *Arthrospira* on fine-scale aspects of individual bee and colony-level health have yet to be characterized.

3 | LIPIDS

While significant research and commercial efforts have been made to match the protein and amino acid contents of pollen, there has been comparatively less focus on the lipid content of PS diets (Ghosh & Jung, 2015; Manning, 2001). Pollen contains a broad diversity of lipids (reviewed in Ischebeck, 2016), that are highly variable depending on plant species (Chakrabarti, Morré, Lucas, Maier, & Sagili, 2019). Microalgae are renowned for their lipid content as it relates to biofuel applications but are also recognized as a rich source of nutritionally relevant lipids, including polyunsaturated fatty acids (PUFAs; Xue et al., 2018) (Ryckebosch et al., 2014). Linoleic and α -linoleic acids are two PUFAs that are essential for honey bees (Arien, Dag, Zarchin, Masci, & Shafir, 2015). Further, dietary PUFA content is correlated with fat body *vitellogenin* (vg) expression (Wegener, Jakop, Schiller, & Müller, 2018). *Chlorella sorokiniana* has lipid contents of 10–20% dry weight and was shown to increase fat deposition in honey bees (Jehlík et al., 2019). However, other species possess even higher lipid contents (dry weight), such as *Botryococcus* (25–75%), *Nanochloropsis* (31–68%), and *Schizochytrium* (50–77%; Demirbas & Demirbas, 2011).

4 | MICRONUTRIENTS

Micronutrients, which include vitamins and minerals, are as important as macronutrients (Gupta, Kening, & Liang, 2008) but have been mostly overlooked in the context of honey bee nutrition. Honey bees obtain micronutrients from pollen as well as “dirty” or turbid water (Butler, 1940). Analysis of bee-collected pollen revealed seasonal fluctuations in mineral levels of potassium, calcium, magnesium and sodium (Bonoan et al., 2017). Algae may also be collected by bees to supplement macronutrition and micronutrition, although this has not been demonstrated beyond anecdotal observations that bees will interact with algae at turbid water sources (Figure 2). Nevertheless, microalgae represent a valuable source of vitamins (tocopherols, ascorbic acid, B vitamins) and minerals (sodium, potassium, calcium, and magnesium; Fabregas & Herrero, 1990).

5 | ANTIOXIDANTS

Nectar and pollen are the honey bee's dietary sources of antioxidants. The antioxidant properties of pollen are attributed to its phenolic content, which may ameliorate cellular damage incurred by oxidative stress. Microalgae are rich in antioxidants, including phenolics (Li et al., 2007) and various photosynthetic pigments including



FIGURE 2 Honey bee at a turbid water source featuring algae at the ground-water interface

carotenoids (Goiris et al., 2012). The microalgal carotenoid astaxanthin is well-studied for its potent antioxidant and immunomodulating properties (Higuera-Ciapara, Felix-Valenzuela, & Goycoolea, 2006). Currently, the main source of microalgal astaxanthin is *Haematococcus pluvialis*, but additional species are under investigation for the production of astaxanthin and related carotenoids (Ambati, Phang, Ravi, & Aswathanarayana, 2014).

6 | PREBIOTIC AND FUNCTIONAL POLYSACCHARIDES

Honey bee guts are populated by a microbial consortium, or microbiota, that is the subject of significant research interest due to its implications in health and disease (Raymann & Moran, 2018). Evidence has emerged demonstrating the beneficial effects of some diet ingredients and bioactive compounds on animal hosts or gut microbiota beyond their primary nutritional roles. Among these “functional ingredients,” the polysaccharides are a promising group (Xu, Xu, Ma, Tang, & Zhang, 2013). This includes dietary fibers, which are broadly defined as non-digestible carbohydrates and lignins found in plants. Some dietary fibers are prebiotic, meaning they may be fermented by the gut microbiota, conferring a broad range of physiological benefits to the host animal (Slavin, 2013). Microalgae exhibit a wide variety of polysaccharide-rich extracellular matrices, which are involved in various physiological processes and environmental interactions (Xiao & Zheng, 2016). Indeed, microalgae have recently been identified as an emergent source of prebiotics with potential to modulate the microbiome, and consequently alter the trajectory of various disease states (de Jesus Raposo, De Morais, & De Morais, 2015). Structurally similar to pollen, microalgal extracellular matrices incorporate cellulose, hemicelluloses (xyloglucan, mannans, glucuronan, β -glucans), and pectin-like carbohydrates (Xiao & Zheng, 2016). In honey bees, indigestible pollen fibers act as fermentation substrates for the core gut microbiota groups *Lactobacillus*, *Bifidobacterium*, and *Gilliamella* (Kešnerová et al., 2017). The major fermentative products produced by the bee microbiota are organic acids, which influence bee metabolism, behavior, (Zheng, Powell, Steele, Dietrich, & Moran, 2017) and immune function (Kwong, Mancenido, & Moran, 2017). In vitro and in vivo models indicate that microalgal biomass can stimulate the abundance and metabolism of *Lactobacillus*, a ubiquitous group of beneficial lactic acid bacteria that are core to many insect and vertebrate microbiota (de Jesus Raposo et al., 2015; Moreno, Corzo, Montilla, Villamiel, & Olano, 2017; Parada, de Caire, de Mulé, & de Cano, 1998). Since bee-associated bacterial symbionts co-evolved with a pollen diet, the gut microbiome phenotype could be a useful metric to evaluate the non-nutritive effects of artificial diets on bee physiology (Ricigliano et al., 2017). Intriguingly, a sulphated polysaccharide from the microalga *Porphyridium* was shown to impact honey bee gut health through a non-prebiotic mechanism (Roussel et al., 2015). In this study, feeding the isolated polysaccharide led to decreased parasite load and bee mortality due to

infection by the gut parasite *Nosema ceranae*, possibly via disruption of *Nosema* adhesion to the gut epithelium. Taken together, the incorporation of microalgal biomass into bee diets has potential for diet-based microbiota manipulations to improve disease resistance.

7 | CONCLUSION AND FUTURE PERSPECTIVE

Malnutrition is increasingly recognized as a major stress factor underlying managed honey bee colony mortality throughout the United States and across the world (López-Urbe, Ricigliano, & Simone-Finstrom, 2020). As a result, there are multiple ongoing efforts to improve bee nutrition. One approach focuses on improving the quality of floral resources through supplemental plantings and landscape enrichment that could augment pollinator nutrition (Decourtye, Mader, & Desneux, 2010). While still early in its implementation, there are limited reports demonstrating a significant impact of supplemental plantings on commercially managed honey bees whereas semi-natural conservation habitats appear to have a stronger positive influence on bee health (Alaux et al., 2017; Ricigliano et al., 2019). Taking into consideration the ecological impacts of climate change and a growing human population, feeding honey bees and other livestock should not compete for arable land that could otherwise be dedicated to conservation efforts or human food production (Draaisma et al., 2013; Lamminen et al., 2019). Intensively cultivated landscapes may not provide the floral diversity necessary for optimal bee nutrition (Donkersley et al., 2014). Microalgae represent an alternative approach to the growing problem of malnutrition, with many species possessing all of the EAAs and lipids required by bees. Similar to current PSs employed by beekeepers, microalgae patties can be applied inside bee hives, above the brood nest (Figure 3). This method is optimal for efficiently feeding the thousands of commercially managed colonies that are involved in agricultural pollination. Microalgae cultivation is highly efficient, done without pesticides or antibiotics and requires minimal water inputs compared to terrestrial plant protein feed resources (Lamminen et al., 2019). Adapting beekeeping and broader livestock management practices with microalgae feeds could contribute to achieving objectives outlined in the United Nations sustainable development goals (SDGs; <https://sustainabledevelopment.un.org/sdgs>) related to food security (SDG 2), sustainable water management (SDG 6), sustainable consumption (SDG 12), climate change (SDG 13), reversal of land degradation, and halting biodiversity loss (SDG 15). In order for this goal to be realized, in depth characterization of the effects of different microalgae species on bee physiology and health are necessary.



FIGURE 3 Delivery of a pollen substitute patty containing *Arthrospira* (spirulina) biomass to a honey bee colony

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CONFLICT OF INTERESTS

The authors declare that there are no conflict of interests.

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