

Review

The impact of microalgae-derived bioactive compounds on gut microbiota modulation

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Microalgae are emerging as a valuable source of bioactive compounds with significant potential to modulate gut microbiota and improve human health. This review explores the roles of microalgal-derived polysaccharides, peptides, polyunsaturated fatty acids, polyphenols, pigments, and micronutrients in shaping intestinal microbial composition and function. These compounds promote beneficial bacteria, inhibit pathogens, enhance immune responses, and regulate key metabolic pathways through the production of short-chain fatty acids and other microbial metabolites. Moreover, microalgae exhibit antioxidant, anti-inflammatory, and antitumor properties that may support the prevention of chronic diseases. However, challenges such as bioavailability, safety, individual microbiota variability, and industrial scalability must be addressed. Despite these limitations, microalgae hold promise as sustainable, multifunctional ingredients for developing functional foods and therapeutic interventions. Future research could focus on clinical validation, optimized extraction methods, and personalized nutrition approaches to fully harness the health benefits of microalgae-derived bioactive compounds in modulating the gut microbiota.

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Introduction

With limited water resources, decreasing arable land, and an ever-growing global population, identifying safe and sustainable food sources to meet increasing demand is one of the 21st century's most urgent challenges [1]. As photosynthetic organisms, microalgae, often seen as 'crops of the future,' can thrive in diverse environments, from aquatic to terrestrial, including wastelands and nonarable land, and they are resilient to climate change [2]. Using light-harvesting, the carbon dioxide (CO₂) can be converted into various compounds, including proteins, lipids, vitamins, antioxidants, and pigments, with health-promoting effects that can be utilized in the development of functional foods [3,4]. Despite these advantages, there are safety considerations, including the potential bioaccumulation of microcystin, toxic proteins, and allergens. Therefore, comprehensive risk assessments are necessary for new microalgae proteins to ensure their safety, in line with EC Regulation 258/97 and EU Recommendation 97/618 EC [5].

Types of microalgae include *Chlorophyta* (green algae), *Bacillariophyta* (diatoms), *Ochrophyta* (Ochrophytes), *Euglenophyta* (Euglenoids), *Cryptophyta* (Cryptomonads), *Haptophyta* (Haptophytes), *Dinophyta* (Dinoflagellates), *Rhodophyta* (red algae), and *Cyanobacteria* (blue-green algae) [6]. Moreover, among the large number of available microalgal species, *Spirulina*, *Chlorella*, and *Phaeodactylum tricornutum* (*P. tricornutum*) have been intensively studied [7]. Besides, the *Spirulina*, due to its characteristics and growth conditions, is estimated to be the most commercialized species in the bioresource market, with an estimated to be valued 2025 global market value of USD 522.6 million and forecast to reach USD 884.6 million by 2032 [8]. Microalgae, abundant in unsaturated fatty acids, proteins, carbohydrates, pigments, and other bioactive compounds, present significant potential as a sustainable food source. Microalgae have the potential to produce various bioactive compounds, including polysaccharides, bioactive peptides, lipids (n-3 long-chain polyunsaturated fatty acids (PUFAs)), vitamins, and pigments (β -carotene, astaxanthin, phycocyanin (PC), and fucoxanthin), with health-promoting effects that can be utilized in the development of functional foods [3,4]. Normally, microalgae bioactive compounds will correspond with some potential health benefits, including enhancing oral, cardiovascular, brain, gastrointestinal, and immune functions, and may help prevent non-communicable diseases like diabetes,

hypercholesterolemia, and obesity [4]. The sustainable development and utilization of microalgae resources might be critical for addressing humanity's energy and nutritional needs.

The gut microbiota is essential to overall health by aiding nutrient metabolism, immune function, and disease prevention, but its disruption can cause inflammation and harmful metabolic effects. [9]. For instance, primary factors like genetics and insulin resistance, along with secondary factors such as viruses, drugs, and poor lifestyle, can all contribute to nonalcoholic fatty liver disease by affecting the gut–liver axis [9]. In this sense, precision nutrition involves customizing the intake of micro-nutrients for human needs, and microalgae can be applied as a promising dietary strategy for managing gut dysbiosis and its related complications [10]. Dysbiosis of the gut microbiota has been implicated in the development of type 2 diabetes, obesity, nonalcoholic liver disease, and other cardiometabolic disorders [11]. Additionally, alterations in gut microbiota are associated with the pathogenesis of various conditions, including cancer, arthritis, autism, depression, anxiety, sleep disorders, HIV, hypertension, and gout [11]. The complex, multilayered cell wall of microalgae, composed of polysaccharides, proteins, lipids, and inorganic salts, poses a significant barrier that must be efficiently, safely, and environmentally breached to extract its high-value components [12]. Most of the bioactive compounds in microalgae are involved in digestive behaviors in the oral-gastrointestinal-tract, and interact with gut microbiota in the colon [12]. In addition, this property of microalgae enables it to resist the digestive process of the intestine and effectively interact with the gut microbiota.

The objective of this review is to comprehensively address the microalgae-derived bioactive compounds and modulation in gut microbiota. Moreover, all the literature referenced refers to microalgae or microalgae-derived bioactive compounds. Particular focus will be given to the *in vitro* or *in vivo* findings to better motivate and understand this review.

Microalgal-derived bioactive compounds and modulation in gut microbiota

Among the effects on the health of gut microbiota, previous research identified the role in fatigue and anti-fatigue [13]. An increase in pathogenic bacteria and a decrease in beneficial bacteria coexist when fatigue is present in both rodents and humans, and changes in gut microbiota were reported after intervention with anti-fatigue foods [13]. Microalgae are recognized as a promising protein source and are rich in bioactive compounds, polysaccharides, bioactive peptides, lipids, polyphenols, pigments, and minerals, with potential human health benefits [4]. Disturbance of the gut

microbiota will lead to the occurrence of diseases, and polysaccharides as prebiotics can improve the disease by regulating the composition of gut microbiota [14]. As for the importance of microalgae bioactive compounds of gut microbiota modulation in health benefits, more details can be seen in Figure 1. Furthermore, the chemical composition of microalgal biomass varies significantly across species and is influenced by growth phase, as well as biotic and abiotic factors such as light, temperature, nutrient availability, and salinity [4].

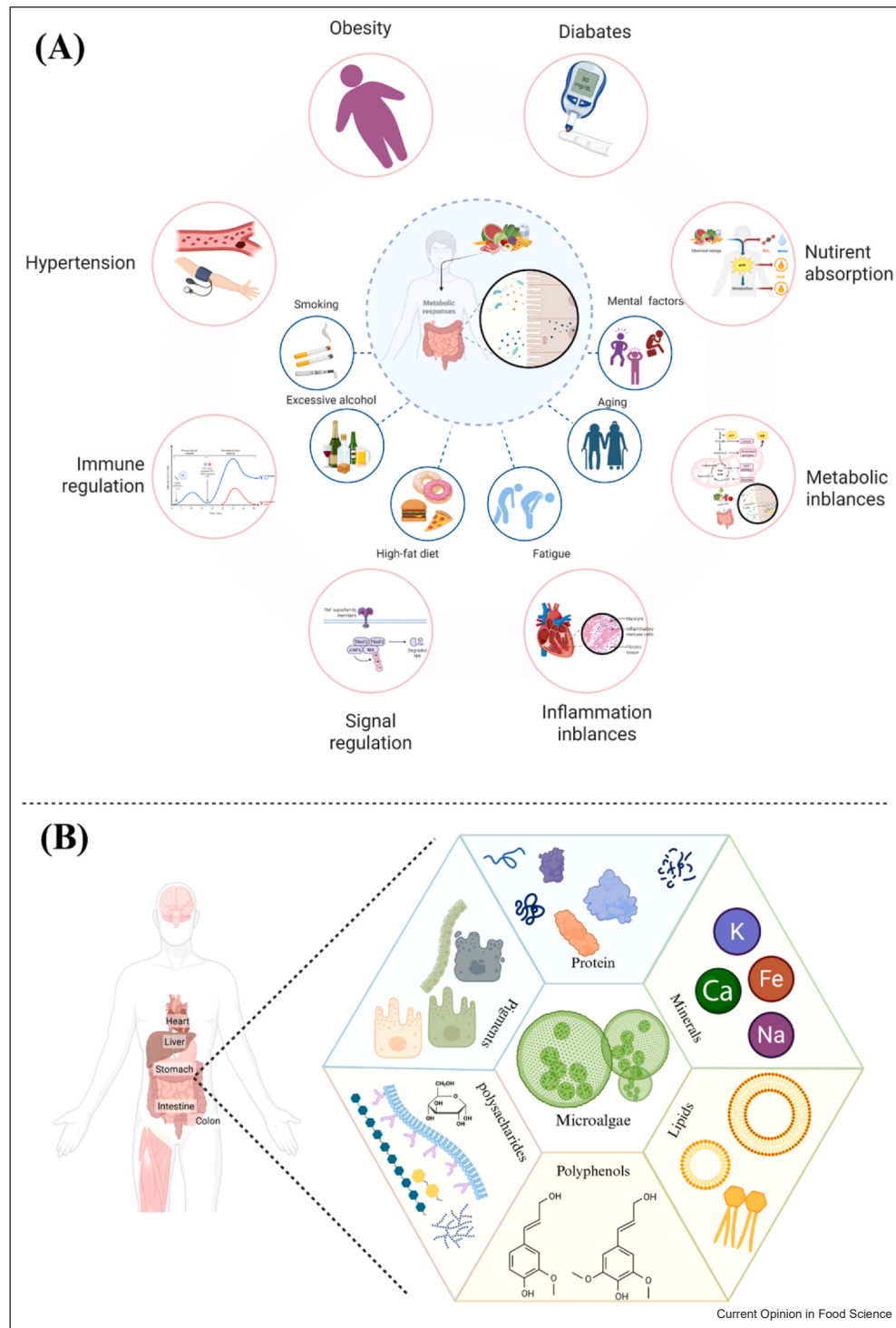
Polysaccharides of microalgae on gut microbiota modulation

Microalgae are rich in carbohydrates, with polysaccharides forming a major component of their cellular structure. These polysaccharides of microalgae fall into three groups [15]: structural cell wall polysaccharides (β -glucans, β -mannans, α -rhamnans, β -galactofuranans, and various heteropolysaccharides), reserve polysaccharides (branched starches and β -glucans), and extracellular polysaccharides (EPS). Microalgal polysaccharides show various biological activities, including immunomodulatory, antitumor, antibacterial, and antioxidant, which can be used as health supplements or can be fermented to produce biofuels [15].

Previous research has comprehensively summarized the interactions between polysaccharides and gut microbiota, highlighting three key aspects [16]: 1) the modulation of gut microbiota composition; 2) the metabolism of polysaccharides into short-chain fatty acids (SCFAs), and 3) the regulation of gut microbiota metabolites, including trimethylamine, tryptophan (Trp), and lipopolysaccharides. The microbiota shares a dynamic relationship with its host, serving vital roles in nutrient metabolism and immune function, and it interacts closely with polysaccharides, which serve as a vital energy source for intestinal microorganisms [14]. Although the host cannot digest polysaccharides, gut microbes break them down, fostering microbial growth and contributing to overall gut health [14]. Microalgae-derived oligosaccharides and phenolics increase the abundance of beneficial gut bacteria, with *Spirulina platensis* and *Desmodesmus maximus* exhibiting the strongest prebiotic effects [17]. Besides, previous research has investigated the polysaccharide obtained from *Spirulina* (PSP) and its effects on the metabolic and gut microbiota of lung cancer in mice [18], revealing that PSP increased the abundance of *Lactobacillus*, *Allobaculum*, *Alloprevotella*, and *Olsenella* and decreased *Bacteroides* and *Acinetobacter*, suggesting that PSP can be applied to against lung cancer diseases. Moreover, the indigestible carbohydrates, including oligosaccharides, fibers, and lactose, could be converted into SCFAs by bacteria residing in the large intestine [11].

As a vital component of the human body, the intestinal microbiota plays a crucial role in maintaining overall

Figure 1



Importance of gut microbiota modulation in health benefits **(a)**. The regulation of the human gut microbiota is influenced by factors such as smoking, excessive alcohol, HFD, fatigue, aging, and mental factors. Additionally, the balance of gut microbiota modulation plays a crucial role in various aspects of human health, including obesity, hypertension, immune regulation, signal regulation, inflammation, metabolic, nutrient absorption, and diabetes. Modulation of gut microbiota by main nutrients and bioactive compounds in microalgae **(b)**. The main nutrients and bioactive compounds in microalgae, such as proteins, minerals, lipids, polyphenols, polysaccharides, and pigments, hold significant potential for modulating human gut microbiota. (Created in Bio Render.com).

health. Investigating the impact of microalgae-derived bioactive compounds on regulating intestinal probiotics could have significant and far-reaching implications.

Bioactive peptides of microalgae on gut microbiota modulation

Microalgae are a promising source of protein, offering a nutrient-dense and consistent source. Microalgae proteins possess functional properties, including solubility, emulsification, foaming, gelling, and natural pigmentation, all of which are produced without competing for arable land [19]. Microalgae-derived proteins exhibit diverse bioactive properties, as evidenced by their demonstrated anti-inflammatory, anticancer, antidiabetic, antibacterial, immune-boosting, neuroprotective, hepatoprotective, and antioxidative effects [20]. Studies indicate that large molecular weight peptides from microalgae are initially broken down in the stomach by pepsin. These larger molecular weight peptides are further broken down in the small intestine by enzymes such as trypsin, chymotrypsin, elastase, carboxypeptidase, and other pancreatic enzymes, ultimately yielding free amino acids (AAs) [12].

One notable AA is Trp, an essential component obtained from dietary protein, which serves as a precursor for various metabolites produced by tissue cells and gut microbiota [21]. These metabolites, including SCFAs, Trp, and bile acid metabolites, play crucial roles in maintaining intestinal barrier integrity, promoting gut homeostasis, and regulating immune function. They also influence genetic, epigenetic, and metabolic pathways in both immunosuppressive and proinflammatory immune cells [22]. Furthermore, dietary protein bioactive peptides can enhance intestinal function by improving the diversity and composition of the gut microbiota, which may help reduce inflammation, oxidative stress, and promote overall host metabolism.

Fatty acids of microalgae on gut microbiota modulation

PUFAs produced by microalgae are natural, high-purity, high-quality fatty acids derived from a sustainable, vegetarian source and offer superior sensory qualities compared to animal-based lipids [23]. PUFAs are characterized by multiple double bonds, being in microalgae mainly omega-3 (C20:5 eicosapentaenoic acid (EPA) and C22:6 docosahexaenoic acid (DHA)) are found in microorganisms, plants, and aquatic animals, and when consumed, algal PUFAs can enhance animals' fatty acid profiles (omega-6/omega-3 ratio) and immune function [24]. Recent studies have highlighted promising algal strains for producing PUFAs, indicating that microalgae could emerge as an innovative source of PUFAs for both human and animal nutrition [25].

Furthermore, microalgae produce a higher amount of omega-3 in contrast to animal-based oils [26].

Specifically, *Isochrysis galbana* and *Phaeodactylum tri-cornutum* yield EPA at levels more than twice as high as those found in traditional sources, while *Cryptocodinium cohnii* produces nearly six times the amount of DHA compared to cod liver oil [26]. Effects of *Spirulina platensis* 55% ethanol extract (SPL55) enriched with PUFAs have the potential to improve serum and hepatic lipid metabolic disorders in rats with hyperlipidemia and can regulate the mRNA and protein involved in lipid metabolism in high-fat diet (HFD)-induced hyperlipidemic rats [27]. Disruptions in gut microbiota can impair lipid and carbohydrate metabolism by negatively affecting liver function, thereby altering enterohepatic circulation and increasing gut permeability. This disruption can alter metabolic pathways such as gluconeogenesis and beta-oxidation, compromising the liver's ability to regulate blood glucose and lipid levels [28].

Polyphenols of microalgae on gut microbiota modulation

Dietary polyphenols play an irreplaceable role in improving and regulating gut microbiota [9]. Microalgae can be characterized as a source of polyphenols, including catechols, gallic acid, tannins, terpenoids, and flavonoids [29]. The development and application of microalgae polyphenols are an emerging trend in functional foods or human health promotion that can be used to prevent diseases associated with fat accumulation inhibition, cardiovascular alleviation, and antioxidation [12]. Polyphenol-enriched foods are closely related to gut microbiota, perhaps owing to their significant antioxidative properties [13].

The effects of digested microalgae biomass of *Chlorella vulgaris*, *Desmodesmus maximus*, *Chlorococcum* sp. cf *hypnosporum*, and *Spirulina platensis* or fructooligosaccharides (FOS) on human colonic microbiota during *in vitro* fermentation (48 h) were evaluated by previous researchers [17]. These results revealed that crude microalgae extraction (containing phenolics and oligosaccharides) could increase gut beneficial bacteria abundance by increasing *Lactobacillus-Enterococcus* and *Bifidobacterium* relative abundances and decreasing *Prevotellaceae-Bacteroidaceae*, *Clostridium histolyticum*, and *Eubacterium rectale-Clostridium coccoides*.

Pigments of microalgae on gut microbiota modulation

Microbial pigments, produced by bacteria, fungi, and microalgae, support intestinal and overall health by regulating gut microbiota and strengthening the intestinal barrier [30]. Their antioxidant, antibacterial, and anti-inflammatory properties promote beneficial microbes, inhibit harmful ones, and may help prevent diseases like obesity, diabetes, and cancer [30]. The major classes of photosynthetic pigments that occur among the microalgae are chlorophylls, carotenoids, and phycobiliproteins [31]. Microalgae, particularly *Spirulina*, are a major source of

PC, a natural blue pigment that is increasingly popular for its health benefits and food-safe colorant properties [5].

Gut microbiota plays a crucial role in regulating the response to immune checkpoint therapy; therefore, modulation of the microbiome with bioactive molecules like carotenoids (α -carotene, β -carotene, lycopene, phytoene, lutein, β -cryptoxanthin, zeaxanthin, astaxanthin, and fucoxanthin) might be a very effective strategy to reduce the risk of chronic diseases [32]. Gut microbiota can convert carotenoids into more easily absorbable forms and generate various bioactive metabolites that benefit human health. For instance, carotenoids are promising molecules that influence gut microbiota by activating immune cells, which may help reduce oxidative stress [32]. The relative researcher has concluded that the natural pigments can regulate the intestinal microbiota, according to [30]: 1) reinforcing the intestinal barrier and immune function by encouraging beneficial microbes and suppressing harmful ones; 2) promoting the bio-transformation and production of active metabolites such as acetylcholine, AAs, SCFAs, and EPS.

A significant portion of microalgae-derived carotenoids reaches the colon, where they interact with gut

microbiota. Some of these carotenoids (β -carotene, astaxanthin, lutein and zeaxanthin, fucoxanthin, lycopene, canthaxanthin, α -carotene, β -cryptoxanthin, phytoene, and siphonaxanthin) can be converted into other compounds during digestion, although the bioavailability of most microalgal carotenoids remains relatively low [33]. Despite their generally low bioavailability, these unabsorbed carotenoids play a vital role in modulating gut microbiota and exhibit significant potential to influence host metabolism [33].

Minerals and vitamins of microalgae on gut microbiota modulation

Vitamins are essential organic micronutrients needed in small amounts for metabolic health, and algae, both seaweed and microalgae, are a natural source [34]. Microalgae can synthesize various vitamins with biotechnological uses, especially β -carotene (a vitamin A precursor), along with water-soluble vitamins B1, B2, B6, B12, and C, and the liposoluble, antioxidant vitamin E [35]. Vitamin content varies by species, and the production, extraction, and purification of these vitamins support antioxidant activity, carbohydrate metabolism, body homeostasis, and the development of functional foods. In terms of some significant bioactive activities of microalgae bioactive

Table 1

Some bioactive activities of microalgae-derived bioactive compounds.

Microalgae-derived bioactive compounds (crude/purified compounds)	Microalgae source	Molecular	Bioactivities	References
Microalgal extracellular polymeric substances	Contain proteins, lipids, exopolysaccharides, and nucleic acids of microalgae	No data	Antiviral capability, inhibition of cancer cell growth, dilatation of bronchial tubes cough relief, lowering blood sugar, antimicrobial capability, loads of drugs, protection against UV rays, moisturizing, and increasing elasticity.	[15]
Oligosaccharide SPO-1	Purified polysaccharides from <i>Spirulina platensis</i>	510.8–997.0 Da	Significantly improved the microbial community by increasing the community richness and diversity; increasing the abundance of more beneficial species (including <i>Bacteroides</i> , <i>Escherichia-Shigella</i> , <i>Megamonas</i> , <i>Megasphaera</i> , <i>Blautia</i> , <i>Bifidobacterium</i> , and <i>Lactobacillus</i>).	[36]
Bioactive peptides	Hydrolysis of protein extracted from microalgae	Protein fragments are composed of 2–20 AAs.	Antioxidant, antidiabetic, antihypertensive, antibacterial, anticancer, osteogenic, antiaging, antiviral.	[37]
Antibacterial peptides (KLENCN YAVELGK)	Pepsin hydrolysate of <i>Limnospira</i> Sp., and then purification of peptides	466.68 Da	Antimicrobial activity against <i>E. coli</i> and <i>S. aureus</i>	[38]
Unsaturated fatty acid, DHA, and EPA	Microalgae lipids	No data	Antibacterial, antifungal, antiinflammatory, alleviating cardiovascular, and modulating gut microbiome.	[39]
β -carotene, astaxanthin, lutein and zeaxanthin, fucoxanthin, lycopene, and others	Microalgae-derived carotenoids	No data	Anti-oxidation, anti-obesity, anticancer, anti-inflammation, cardiovascular health, vision protection, skin protection, and others.	[33]
β -carotene (pigments)	Crude extraction or purification from microalgae	No data	anti-oxidation (promote lifespan in lab animals); anti-inflammation (reduce oxidative stress in culture cells)	[40]

compounds and some related research, more details can be observed in Table 1 [15,33,36–40].

Microalgae are a rich reservoir of vitamins (B vitamins and vitamin D) and minerals (including essential minerals such as iron (Fe), calcium (Ca), and magnesium (Mg)) [23]. Minerals constitute a significant fraction of microalgal biomass, accounting for 2–40% of its dry weight [41]. Major minerals include Ca, Mg, phosphorus (P), potassium (K), and sodium (Na), while trace minerals consist of manganese (Mn), Fe, selenium (Se), zinc (Zn), and copper (Cu) [41]. These minerals are vital for numerous biological processes [42]. However, excessive mineral supplementation may negatively affect intestinal integrity and increase susceptibility to enteric infections, potentially by altering gut microbiota composition and function [43].

In vitro selenium (Se) bioaccessibility combined with *in vivo* biostability and bioactivity in Se-enriched microalgae (*Chlorella sorokiniana*) has been evaluated [44]. In that study, the results revealed that Se-enriched microalgae were 81% *in vitro* gastrointestinal digestion, whereas the *in vivo* mice supplemented with Se-enriched *Chlorella sorokiniana* showed a high concentration of selenium in the kidney, reflecting a potential mechanism of excretion by urine. Additional studies examining the effects of microalgal biomass on gut microbiota modulation, both *in vitro* and *in vivo*, are summarized in Tables 2 [44–47] and 3 [7,17,48,49].

Challenges and considerations

While microalgal-derived bioactive compounds present promising opportunities for gut microbiota modulation, several challenges and considerations must be addressed before their widespread application. One major challenge is bioavailability [50], such as polyphenols and carotenoids exhibit low absorption rates and limited stability in the gastrointestinal tract, which may reduce their effectiveness. Additionally, the complexity and variability of individual gut microbiota can lead to inconsistent responses among different populations, making it difficult to generalize therapeutic outcomes. Another critical concern is safety. Certain microalgae may accumulate harmful substances such as heavy metals, allergens, or toxins like microcystins, necessitating rigorous screening, purification, and regulatory oversight. Standardization of extraction and processing methods also poses a hurdle; variations in cultivation conditions, strain selection, and extraction techniques can significantly influence the quality and efficacy of bioactive compounds.

Moreover, economic and technological barriers remain, as large-scale production and downstream processing of high-purity compounds are often cost-intensive and technically demanding. Lastly, there is a lack of long-term clinical studies to support health claims and establish dosage guidelines for different populations. To realize the full potential of microalgal bioactives, comprehensive multidisciplinary research integrating

Table 2

Studies on gut microbiota modulation effects *in vivo* assays with microalgae-derived bioactive compounds.

Microalgae	Microalgae biomass	Assay	Major outcomes	References
<i>Spirulina platensis</i>	polysaccharides	<i>In vivo</i>	<i>Spirulina platensis</i> polysaccharides (SPLP) had significant hypolipidemic and hypoglycemic effects in HFD-fed rats; SPLP significantly modulated the intestinal microbiota dysbiosis.	[45]
<i>Scenedesmus obliquus</i>	protein, carotenoids,	<i>In vivo</i> with Wistar rats	High values of protein efficiency ratio after the consumption of microalga; ingestion of microalga <i>S. obliquus</i> reduced serum glucose of up to 42%; the intake of the microalga reduced triglycerides of up to 70%; the atherogenic index decreased with increasing consumption of <i>S. obliquus</i> .	[46]
<i>Spirulina</i>	lipids	<i>In vivo</i>	Dietary <i>Spirulina</i> lipids (SL) reduced the body weight in diet-induced obese mice; SL supplementation also suppressed the hepatic lipid accumulation in obese mice; the anti-obesity and antifatty liver effects of SL were comparable with fish oil; possible mechanisms might be through regulating SREBP-1c, SREBP-2, and PPAR α .	[47]
<i>Chlorella sorokiniana</i> (selenium (Se)-enriched microalgae)	Selenium (minerals)	<i>In vivo</i>	The Se bioaccessibility in Se-enriched <i>Chlorella sorokiniana</i> evaluated by <i>in vitro</i> gastrointestinal digestion of the selenized microalga was 81% (79% as SeMet); mice supplemented with Se-enriched <i>C. sorokiniana</i> presented increased Se concentration in the kidney; Se bioavailability and bioactivity were 76–85% in serum from these mice animals	[44]

Table 3

Studies on gut microbiota modulation effects *in vitro* assays with microalgae-derived bioactive compounds.

Microalgae	Microalgae biomass	Assay	Major outcomes	References
<i>Caulerpa lentillifera</i> J. Agardh, <i>Ulva rigida</i> C. Agardh, <i>Spirulina</i> sp., <i>Coelastrella</i> sp. (KKUP1), and <i>Scenedesmus</i> sp. (KKUP2)	Protein	<i>In vitro</i> digestibility	<i>Spirulina</i> sp. had the highest protein content (44.44% of dry matter basis; $P < 0.0001$) and was rich in AAs (Lys, Thr, and Trp). Moreover, protein solubility, digestibility (DPD and GPD _{Protein}), and digestible protein of <i>Spirulina</i> sp. were significantly higher than those of other species (52.35%, 49–67% and 20.70%, respectively). Energy digestibility (GPD _{Energy}) and digestible energy were highest in <i>Ulva rigida</i> (99.40% and 15.29 MJ kg ⁻¹ , respectively; $P < 0.0001$).	[48]
<i>Chlorella vulgaris</i> (CHL), <i>Desmodesmus maximus</i> (DES), <i>Chlorococcum</i> sp. cf <i>hypnosporum</i> (CHLO), and <i>Spirulina platensis</i> (SPI)	Oligosaccharides and phenolics	<i>In vitro</i> (in vitro colonic fermentation of digested)	Increase gut beneficial bacteria abundance by increasing <i>Lactobacillus-Enterococcus</i> (similar increase in the relative abundance for DES, SPI, and FOS, approximately 30%) and <i>Bifidobacterium</i> relative abundances (similar increase in the relative abundance in CHL and SPI, approximately 36–37%).	[17]
<i>Spirulina platensis</i>	PC (pigments)	<i>In vitro</i>	Reduces liver fat accumulation in methionine-choline-deficient diet-induced nonalcoholic steatohepatitis mice; regulates liver lipid metabolism by activating the AMP-activated protein kinase pathway.	[49]
<i>Spirulina</i> , <i>Chlorella</i> , and <i>Phaeodactylum tricornutum</i> (P. <i>tricornutum</i>)	SCFAs	<i>In vitro</i> (in vitro simulated digestion and in vitro colonic fermentation)	The bacterial copy numbers were affected by the <i>in vitro</i> digestion process and colonic fermentation time in the analyses of quantitative polymerase chain reaction (PCR) technology; microalgal biomolecules' digestion promoted the release of SCFAs during <i>in vitro</i> colonic microbiota fermentation, particularly acetic, butanoic, and propanoic, indicating a potential health benefit for the human gut.	[7]

microbiology, nutrition, toxicology, and bioengineering is essential.

Conclusion and future perspectives

In conclusion, microalgae-derived bioactive compounds offer substantial potential for modulating gut microbiota and improving human health. Through mechanisms involving microbial composition shifts, metabolite production (e.g. SCFAs, Trp derivatives), and immune modulation, these compounds influence a wide range of physiological processes. Polysaccharides, peptides, PUFAs, polyphenols, pigments, and essential micro-nutrients derived from microalgae exhibit synergistic effects that promote the growth of beneficial bacteria, suppress pathogens, enhance intestinal barrier integrity, and regulate host metabolism and immune response. Such multifactorial actions position microalgae as promising candidates for next-generation functional foods and therapeutic agents in managing gut-related diseases.

Despite these promising outcomes, significant challenges remain. Bioavailability, compound stability, inter-individual microbiota variability, and long-term effects require further investigation. Moreover, standardized protocols for extraction, purification, and safety assessment are critical to ensure consistency and efficacy across applications. Future research should focus on well-controlled clinical trials to validate *in vivo* outcomes, particularly in patient populations with chronic diseases or metabolic syndromes. Additionally, advancements in bioprocessing technologies could enhance the scalability and economic feasibility of microalgal products.

Looking ahead, integrating microalgal bioactive compounds into personalized nutrition strategies could revolutionize preventive healthcare. A deeper understanding of host–microbiota interactions, aided by omics technologies, will pave the way for tailoring microalgal-based interventions to individual microbiomes,

maximizing therapeutic outcomes while supporting environmental sustainability through algae-based innovation.

Data Availability

The authors are unable or have chosen not to specify which data has been used.

Declaration of Competing Interest

The authors have no conflict of interest to declare.

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