

Editorial: socio-ecology of microbes in a changing ocean

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Editorial on the Research Topic

[Socio-Ecology of Microbes in a Changing Ocean](#)

Marine microbes live in complex socio-ecological networks with diverse microbial and macrobial neighbors. These networks operate on different scales, spanning the single-cell level to microhabitats and (meta)communities to entire phytoplankton blooms ([Faust and Raes, 2012](#) ; [Teeling et al., 2012](#) ; [Cordero and Datta, 2016](#)). Boosted by recent advances in next-generation sequencing and high-resolution chemical analyses, the scientific community begins to unveil the complexity of microbial relationships and how these interactions influence biological dynamics and holobiont functioning ([Pita et al., 2018](#) ; [Van De Water et al., 2018](#)). Ultimately, the elucidation of microbial network processes helps with the interpretation of large-scale ecological and biogeochemical events ([Strom, 2008](#) ; [Fuhrman et al., 2015](#)).

The Frontiers Research Topic Socio-Ecology of Microbes in a Changing Ocean invited contributions on microbial signaling and communication, the effect of microbial interactions on microhabitat structuring, the function of secondary metabolites in bacterial antagonism and microbe-host interactions, the conversion and cross-feeding of nutrients, microbial physiology and gene regulation in response to co-occurring organisms, as well as metabolic exchange in diverse communities. Special emphasis was placed on socio-ecological dynamics under a changing climate, addressing the biological effects of ocean acidification and warming and the increasing spread of invasive species.

The Research Topic encompasses 16 papers on diverse aspects of microbial cooperation and competition, such as bacterial interactions with phytoplankton, macroalgae and invertebrates, chemical microdiversity, interkingdom signaling, viral infections of macroalgae, as well as the distribution of archaea and fungi in relation to environmental parameters. In addition, one Review and one Perspective article offer exciting views on coral disease and phytoplankton multicellularity.

Bacteria-Algae Interactions

Interactions between bacteria and algae are a major focus of the Research Topic, reflecting the pivotal role of micro- and macroalgae in marine carbon fluxes ([Falkowski and Woodhead, 1992](#) ; [Field et al., 1998](#)) and associated microbial dynamics ([Amin et al., 2012](#) ; [Martin et al., 2014](#) ; [Seymour et al., 2017](#)). Contributed papers highlight the importance of bacteria for algae and vice versa, describing the composition and function of associated microbiota, interdependencies by metabolite exchange, and how these interactions can vary under changing environmental conditions.

In their paper “ Bacterial Communities of Diatoms Display Strong Conservation Across Strains and Time,” [Behringer et al.](#) show that the diatoms *Asterionellopsis glacialis* and *Nitzschia longissima* harbor conserved microbiota that remain stable over 1 year of co-existence. These distinct and consistent temporal associations are linked to a range of bacterial processes that enhance diatom growth. The concept of specific associations between bacteria and microalgae is supported by [Crenn et al.](#) in their paper “ Bacterial Epibiotic Communities of Ubiquitous and Abundant Marine Diatoms

Are Distinct in Short- and Long-Term Associations.” Here, screening of single *Thalassiosira* and *Chaetoceros* cells demonstrates that laboratory experiments select for specific diatom microbiota adapted to long-term associations, whereas environmental diatoms harbor different bacterial associates. A step toward deciphering the molecular mechanisms behind such interactions is made by [Torres-Monroy and Ullrich](#) in their paper “ Identification of Bacterial Genes Expressed During Diatom-Bacteria Interactions Using an *in vivo* Expression Technology Approach.” Here, bacterial gene expression is studied in response to the diatom *Thalassiosira weissflogii*, revealing specific expression of bacterial promoters during interactions with *T. weissflogii*. These observations correspond to specific regulation of bacterial attachment, nitrogen metabolism and heavy metal resistance. Interactions between algae and bacteria also include the exchange of ecologically relevant metabolites, as demonstrated by [Wienhausen et al.](#) in “ The Exometabolome of Two Model Strains of the *Roseobacter* Group: A Marketplace of Microbial Metabolites.” Here, ultrahigh-resolution mass spectrometry reveals diverse exometabolites secreted by the bacteria *Phaeobacter inhibens* and *Dinoroseobacter shibae*, including plant auxins and precursors of different B vitamins that may benefit co-occurring auxotrophs. A related scenario is presented in “ The B-Vitamin Mutualism Between the Dinoflagellate *Lingulodinium polyedrum* and the Bacterium *Dinoroseobacter shibae* ” by [Cruz-López et al.](#) who describe that *D. shibae* is dependent on vitamin B₇ produced by *L. polyedrum* while in turn providing vitamins B₁ and B₁₂ to the eukaryotic partner. [Yarimizu et al.](#) describe another type of bacterial interaction with this dinoflagellate species

in their paper “ Iron and Harmful Algae Blooms: Potential Algal-Bacterial Mutualism Between *Lingulodinium polyedrum* and *Marinobacter algicola* .” Here, *M. algicola* is shown as essential for dinoflagellate growth by supplying bioavailable iron via the siderophore vibrioferrin. As *L. polyedrum* can be a major cause of harmful algal blooms, bacteria-algae interactions can hence also have detrimental ecological and economic consequences. This alternative perspective is also addressed by [Bramucci et al.](#) in “ The Bacterial Symbiont *Phaeobacter inhibens* Shapes the Life History of Its Algal Host *Emiliana huxleyi* .” During long-term co-cultivation, *P. inhibens* selectively kills calcifying and flagellated types of the coccolithophore *Emiliana huxleyi* , whereas non-calcifying *E. huxleyi* remain unaffected. This differential pathogenesis may alter the composition of *E. huxleyi* blooms, with probable consequences for marine primary production.

The section on bacteria-algae interactions also includes two papers on macroalgae and associated microbes. Macroalgae are important primary producers in coastal environments and have central functions as habitat formers and nutrient source, but are threatened by environmental stressors ([Steneck and Erlandson, 2002](#) ; [Krumhansl et al., 2016](#)). In their paper “ Novel ssDNA Viruses Detected in the Virome of Bleached, Habitat-Forming Kelp *Ecklonia radiata* ,” [Beattie et al.](#) shotgun-sequenced viral particles isolated from healthy and diseased phenotypes of the kelp *Ecklonia radiata* . This approach identified novel ssDNA viruses restricted to bleached kelp, indicating that stress-induced viral infections may affect coastal primary production. The view on macroalgae-associated microbes is complemented by the paper “ Exploring the Cultivable *Ectocarpus* Microbiome,” where <https://assignbuster.com/editorial-socio-ecology-of-microbes-in-a-changing-ocean/>

[KleinJan et al.](#) isolated over 300 bacterial strains associated with the brown macroalga *Ectocarpus subulatus*. This first step toward a model system for functional studies of algae–bacteria interactions during abiotic stress is highly relevant for future-ocean scenarios in the wake of climate change.

The collection of papers related to bacteria and algae concludes with a Perspective article raising thought-provoking concepts about “Multicellular Features of Phytoplankton.” Here, [Abada and Segev](#) propose that microalgal populations often display the characteristics of a multicellular-like community; representing an evolutionary intermediate between single cells and aggregates that communicate and cooperate. By combining evidence on coccolithophores and diatoms, two key phytoplankton groups, the authors discuss exciting aspects such as coordinated behavior and programmed cell death in a multicellular context.

Bacteria-Cnidaria Interactions

Studies on functional interactions between cnidarian hosts and microbes are receiving continued interest, particularly owing to ecological threats connected to climate change, for instance coral bleaching ([Pandolfi et al., 2003](#) ; [Bourne et al., 2016](#)). This Research Topic includes four related papers, including the Review “Responses of Coral-Associated Bacterial Communities to Local and Global Stressors.” Here, [McDevitt-Irvin et al.](#) summarize 45 recent studies to show that coral health can be strongly influenced by microbiome composition, illustrating that stress-related shifts in bacterial diversity may have important ecological consequences. A second coral-related contribution, “Quorum Sensing Interference and Structural

Variation of Quorum Sensing Mimics in Australian Soft Coral” by [Freckelton et al.](#) describes how coral-derived metabolites mimic bacterial signaling molecules and hence influence cell-cell communication. The finding of chemical crosstalk between soft corals and their associated bacteria is important considering the ecological network within the coral holobiont. Another type of cnidarian, the sea anemone *Nematostella*, is investigated in the paper “ Predicted Bacterial Interactions Affect *in vivo* Microbial Colonization Dynamics in *Nematostella*” by [Domin et al.](#) Through bacterial cultivation and co-occurrence networks, bacteria-bacteria interactions are shown to change according to the host's developmental stage. Predicted competitive bacteria influence community structure for a short period of time but are soon replaced, indicating a high degree of resilience within the bacterial community. In their paper “ Stimulated Respiration and Net Photosynthesis in *Cassiopeia* sp. during Glucose Enrichment Suggests *in hospite* CO₂ Limitation of Algal Endosymbionts,” [Rädecker et al.](#) show that glucose enrichment stimulates respiration and photosynthesis in the holobiont of upside-down jellyfish, likely resulting from bacterial activity that subsequently stimulates primary production by algal symbionts through increased CO₂ availability.

Effects of Environmental Parameters on Microbial Distribution

The third section of the Research Topic includes two papers on the distribution of microbes in relation to environmental parameters ([Hanson et al., 2012](#) ; [Sunagawa et al., 2015](#)). Whereas biogeographical patterns are comparatively well-studied for bacteria, [Müller et al.](#) focus on archaea by

resolving “Spatiotemporal Dynamics of Ammonia-Oxidizing Thaumarchaeota in Distinct Arctic Water Masses.” The authors describe distributional patterns of Thaumarchaeota genotypes in specific water masses in the Arctic-Atlantic boundary, a region under special threat from climate change. Considering the thaumarchaeotal contribution to ammonia oxidation, these findings have implications for nitrogen cycling. Finally, the paper “Spatiotemporal Distribution and Assemblages of Planktonic Fungi in the Coastal Waters of the Bohai Sea” by [Wang et al.](#) illustrates regional and temporal changes in the abundance and diversity of fungi, a so-far largely unexplored group of marine microbes. Differential distribution of *Ascomycota* and *Basidiomycota* between coastal habitats was related to riverine inputs and phytoplankton detritus, providing insights into the ecology of fungi in marine systems.

Conclusions and Perspectives

As evidenced by the contributions to this Research Topic, the socio-ecology of microbes is a growing field of research that substantially benefits from cross-disciplinary approaches. The diverse Topic contributions showcase that the field is clearly moving from merely descriptive analyses of “who-is-associated-with-whom” to functional studies of diversity and phenotypic traits within metaorganisms, which harbor a network of associates from all domains of life. Holistic perspectives and systemic analyses, which acknowledge the complexity and linkages within biological systems, are fundamental for the comprehensive understanding of ecological and biogeochemical processes, particularly in the wake of climate change.

Author Contributions

This editorial was co-authored by the Topic Editors MW, SL, and TH.

Conflict of Interest Statement

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.

References

Amin, S. A., Parker, M. S., and Armbrust, E. V. (2012). Interactions between diatoms and bacteria. *Microbiol. Mol. Biol. Rev.* 76, 667–684. doi: 10.1128/MMBR.00007-12

[PubMed Abstract](#) | [CrossRef Full Text](#) | [Google Scholar](#)

Bourne, D. G., Morrow, K. M., and Webster, N. S. (2016). Insights into the coral microbiome: underpinning the health and resilience of reef ecosystems. *Annu. Rev. Microbiol.* 70, 317–340. doi: 10.1146/annurev-micro-102215-095440

[PubMed Abstract](#) | [CrossRef Full Text](#) | [Google Scholar](#)

Cordero, O. X., and Datta, M. S. (2016). Microbial interactions and community assembly at microscales. *Curr. Opin. Microbiol.* 31, 227–234. doi: 10.1016/j.mib.2016.03.015

[PubMed Abstract](#) | [CrossRef Full Text](#) | [Google Scholar](#)

Falkowski, P. G., and Woodhead, A. D. (1992). *Primary Productivity and Biogeochemical Cycles in the Sea*. New York, NY: Plenum Press.

[Google Scholar](#)

Faust, K., and Raes, J. (2012). Microbial interactions: from networks to models. *Nat. Rev. Microbiol.* 10, 538–550. doi: 10. 1038/nrmicro2832

[PubMed Abstract](#) | [CrossRef Full Text](#) | [Google Scholar](#)

Field, C. B., Behrenfeld, M. J., Randerson, J. T., and Falkowski, P. (1998). Primary production of the biosphere: integrating terrestrial and oceanic components. *Science* 281, 237–240. doi: 10. 1126/science. 281. 5374. 237

[PubMed Abstract](#) | [CrossRef Full Text](#) | [Google Scholar](#)

Fuhrman, J. A., Cram, J. A., and Needham, D. M. (2015). Marine microbial community dynamics and their ecological interpretation. *Nat. Rev. Microbiol.* 13, 133–146. doi: 10. 1038/nrmicro3417

[PubMed Abstract](#) | [CrossRef Full Text](#) | [Google Scholar](#)

Hanson, C. A., Fuhrman, J. A., Horner-Devine, M. C., and Martiny, J. B. H. (2012). Beyond biogeographic patterns: processes shaping the microbial landscape. *Nat. Rev. Microbiol.* 10, 497–507. doi: 10. 1038/nrmicro2795

[PubMed Abstract](#) | [CrossRef Full Text](#) | [Google Scholar](#)

Krumhansl, K. A., Okamoto, D. K., Rassweiler, A., Novak, M., Bolton, J. J., Cavanaugh, K. C., et al. (2016). Global patterns of kelp forest change over

<https://assignbuster.com/editorial-socio-ecology-of-microbes-in-a-changing-ocean/>

the past half-century. *Proc. Natl. Acad. Sci. U. S. A.* 113, 13785–13790. doi: 10.1073/pnas.1606102113

[PubMed Abstract](#) | [CrossRef Full Text](#) | [Google Scholar](#)

Martin, M., Portetelle, D., Michel, G., and Vandenberg, M. (2014). Microorganisms living on macroalgae: diversity, interactions, and biotechnological applications. *Appl. Microbiol. Biotechnol.* 98, 2917–2935. doi: 10.1007/s00253-014-5557-2

[PubMed Abstract](#) | [CrossRef Full Text](#) | [Google Scholar](#)

Pandolfi, J. M., Bradbury, R. H., Sala, E., Hughes, T. P., Bjorndal, K. A., Cooke, R. G., et al. (2003). Global trajectories of the long-term decline of coral reef ecosystems. *Science* 301, 955–958. doi: 10.1126/science.1085706

[PubMed Abstract](#) | [CrossRef Full Text](#) | [Google Scholar](#)

Pita, L., Rix, L., Slaby, B. M., Franke, A., and Hentschel, U. (2018). The sponge holobiont in a changing ocean: from microbes to ecosystems. *Microbiome* 6: 46. doi: 10.1186/s40168-018-0428-1

[PubMed Abstract](#) | [CrossRef Full Text](#) | [Google Scholar](#)

Seymour, J. R., Amin, S. A., Raina, J. B., and Stocker, R. (2017). Zooming in on the phycosphere: the ecological interface for phytoplankton-bacteria relationships. *Nat. Microbiol.* 2, 17065. doi: 10.1038/nmicrobiol.2017.65

[PubMed Abstract](#) | [CrossRef Full Text](#) | [Google Scholar](#)

Steneck, R., and Erlandson, J. M. (2002). Kelp forest ecosystems: biodiversity, stability, resilience and future. *Environ. Conserv.* 29, 436–459. doi: 10.1017/S0376892902000322

[CrossRef Full Text](#) | [Google Scholar](#)

Strom, S. L. (2008). Microbial ecology of ocean biogeochemistry: a community perspective. *Science* 320, 1043–1045. doi: 10.1126/science.1153527

[PubMed Abstract](#) | [CrossRef Full Text](#) | [Google Scholar](#)

Sunagawa, S., Coelho, L. P., Chaffron, S., Kultima, J. R., Labadie, K., Salazar, G., et al. (2015). Structure and function of the global ocean microbiome. *Science* 348: 1261359. doi: 10.1126/science.1261359

[PubMed Abstract](#) | [CrossRef Full Text](#) | [Google Scholar](#)

Teeling, H., Fuchs, B. M., Becher, D., Klockow, C., Gardebrecht, A., Bennke, C. M., et al. (2012). Substrate-controlled succession of marine bacterioplankton populations induced by a phytoplankton bloom. *Science* 336, 608–611. doi: 10.1126/science.1218344

[PubMed Abstract](#) | [CrossRef Full Text](#) | [Google Scholar](#)

Van De Water, J. A. J. M., Allemand, D., and Ferrier-Pagès, C. (2018). Host-microbe interactions in octocoral holobionts - recent advances and perspectives. *Microbiome* 6: 64. doi: 10.1186/s40168-018-0431-6

[PubMed Abstract](#) | [CrossRef Full Text](#) | [Google Scholar](#)

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