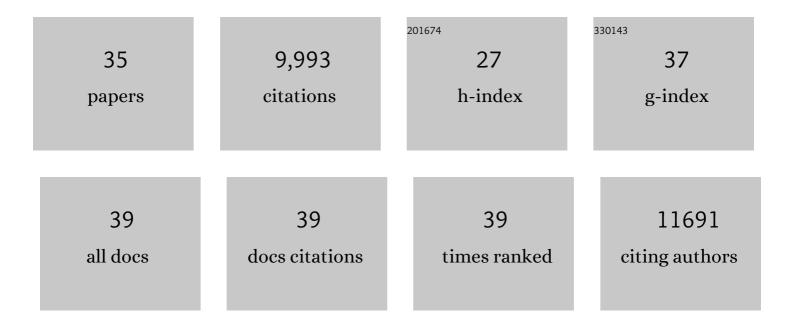
Lucie Bittner

List of Publications by Year in descending order

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LUCIE RITTNER

#	Article	lF	CITATIONS
1	Functional traitâ€based approaches as a common framework for aquatic ecologists. Limnology and Oceanography, 2021, 66, 965-994.	3.1	99
2	Towards omics-based predictions of planktonic functional composition from environmental data. Nature Communications, 2021, 12, 4361.	12.8	16
3	A resource-frugal probabilistic dictionary and applications in bioinformatics. Discrete Applied Mathematics, 2020, 274, 92-102.	0.9	13
4	Community‣evel Responses to Iron Availability in Open Ocean Plankton Ecosystems. Global Biogeochemical Cycles, 2019, 33, 391-419.	4.9	76
5	Mixotrophic protists display contrasted biogeographies in the global ocean. ISME Journal, 2019, 13, 1072-1083.	9.8	55
6	Analysis of the genomic basis of functional diversity in dinoflagellates using a transcriptomeâ€based sequence similarity network. Molecular Ecology, 2018, 27, 2365-2380.	3.9	12
7	The epibiotic life of the cosmopolitan diatom <i>Fragilariopsis doliolus</i> on heterotrophic ciliates in the open ocean. ISME Journal, 2018, 12, 1094-1108.	9.8	26
8	Influence of diatom diversity on the ocean biological carbon pump. Nature Geoscience, 2018, 11, 27-37.	12.9	451
9	Clade-specific diversification dynamics of marine diatoms since the Jurassic. Nature Ecology and Evolution, 2018, 2, 1715-1723.	7.8	40
10	A de novo approach to disentangle partner identity and function in holobiont systems. Microbiome, 2018, 6, 105.	11.1	19
11	A communal catalogue reveals Earth's multiscale microbial diversity. Nature, 2017, 551, 457-463.	27.8	1,942
12	The evolution of diatoms and their biogeochemical functions. Philosophical Transactions of the Royal Society B: Biological Sciences, 2017, 372, 20160397.	4.0	134
13	Acclimation of a low iron adapted Ostreococcus strain to iron limitation through cell biomass lowering. Scientific Reports, 2017, 7, 327.	3.3	25
14	Mare Incognitum: A Glimpse into Future Plankton Diversity and Ecology Research. Frontiers in Marine Science, 2017, 4, .	2.5	10
15	Insights into global diatom distribution and diversity in the world's ocean. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, E1516-25.	7.1	561
16	Benthic protists: the under-charted majority. FEMS Microbiology Ecology, 2016, 92, fiw120.	2.7	94
17	Plankton networks driving carbon export in the oligotrophic ocean. Nature, 2016, 532, 465-470.	27.8	670
18	Marine protist diversity in <scp>E</scp> uropean coastal waters and sediments as revealed by highâ€ŧhroughput sequencing. Environmental Microbiology, 2015, 17, 4035-4049.	3.8	384

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19	Determinants of community structure in the global plankton interactome. Science, 2015, 348, 1262073.	12.6	842
20	Eukaryotic plankton diversity in the sunlit ocean. Science, 2015, 348, 1261605.	12.6	1,551
21	Environmental characteristics of Agulhas rings affect interocean plankton transport. Science, 2015, 348, 1261447.	12.6	158
22	Seasonal diversity and dynamics of haptophytes in the <scp>S</scp> kagerrak, <scp>N</scp> orway, explored by highâ€throughput sequencing. Molecular Ecology, 2015, 24, 3026-3042.	3.9	90
23	Testing ecological theories with sequence similarity networks: marine ciliates exhibit similar geographic dispersal patterns as multicellular organisms. BMC Biology, 2015, 13, 16.	3.8	42
24	Patterns of Rare and Abundant Marine Microbial Eukaryotes. Current Biology, 2014, 24, 813-821.	3.9	450
25	Diversity patterns of uncultured Haptophytes unravelled by pyrosequencing in Naples Bay. Molecular Ecology, 2013, 22, 87-101.	3.9	70
26	Species Diversity, Phylogeny and Large Scale Biogeographic Patterns of the Genus <i>Padina</i> (Phaeophyceae, Dictyotales). Journal of Phycology, 2013, 49, 130-142.	2.3	53
27	454 Pyrosequencing to Describe Microbial Eukaryotic Community Composition, Diversity and Relative Abundance: A Test for Marine Haptophytes. PLoS ONE, 2013, 8, e74371.	2.5	118
28	Effect of elevated CO ₂ on the dynamics of particle-attached and free-living bacterioplankton communities in an Arctic fjord. Biogeosciences, 2013, 10, 181-191.	3.3	26
29	Ocean acidification shows negligible impacts on high-latitude bacterial community structure in coastal pelagic mesocosms. Biogeosciences, 2013, 10, 555-566.	3.3	60
30	The Protist Ribosomal Reference database (PR2): a catalog of unicellular eukaryote Small Sub-Unit rRNA sequences with curated taxonomy. Nucleic Acids Research, 2012, 41, D597-D604.	14.5	1,463
31	An original mode of symbiosis in open ocean plankton. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 18000-18005.	7.1	126
32	Evolutionary history of the Corallinales (Corallinophycidae, Rhodophyta) inferred from nuclear, plastidial and mitochondrial genomes. Molecular Phylogenetics and Evolution, 2011, 61, 697-713.	2.7	119
33	Some considerations for analyzing biodiversity using integrative metagenomics and gene networks. Biology Direct, 2010, 5, 47.	4.6	50
34	Multigene phylogenetic analyses support recognition of the Sporolithales ord. nov Molecular Phylogenetics and Evolution, 2010, 54, 302-305.	2.7	77
35	Molecular phylogeny of the Dictyotales and their position within the Phaeophyceae, based on nuclear, plastid and mitochondrial DNA sequence data. Molecular Phylogenetics and Evolution, 2008, 49, 211-226.	2.7	69