Lucie Bittner

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1577022/publications.pdf

Version: 2024-02-01

201674 330143 9,993 35 27 37 h-index citations g-index papers 39 39 39 11691 docs citations times ranked citing authors all docs

| # | Article | IF | CITATIONS |
|----|---|------|-----------|
| 1 | A communal catalogue reveals Earth's multiscale microbial diversity. Nature, 2017, 551, 457-463. | 27.8 | 1,942 |
| 2 | Eukaryotic plankton diversity in the sunlit ocean. Science, 2015, 348, 1261605. | 12.6 | 1,551 |
| 3 | 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 |
| 4 | Determinants of community structure in the global plankton interactome. Science, 2015, 348, 1262073. | 12.6 | 842 |
| 5 | Plankton networks driving carbon export in the oligotrophic ocean. Nature, 2016, 532, 465-470. | 27.8 | 670 |
| 6 | 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 |
| 7 | Influence of diatom diversity on the ocean biological carbon pump. Nature Geoscience, 2018, 11, 27-37. | 12.9 | 451 |
| 8 | Patterns of Rare and Abundant Marine Microbial Eukaryotes. Current Biology, 2014, 24, 813-821. | 3.9 | 450 |
| 9 | Marine protist diversity in <scp>E</scp> uropean coastal waters and sediments as revealed by highâ€throughput sequencing. Environmental Microbiology, 2015, 17, 4035-4049. | 3.8 | 384 |
| 10 | Environmental characteristics of Agulhas rings affect interocean plankton transport. Science, 2015, 348, 1261447. | 12.6 | 158 |
| 11 | The evolution of diatoms and their biogeochemical functions. Philosophical Transactions of the Royal Society B: Biological Sciences, 2017, 372, 20160397. | 4.0 | 134 |
| 12 | 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 |
| 13 | 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 |
| 14 | 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 |
| 15 | Functional traitâ€based approaches as a common framework for aquatic ecologists. Limnology and Oceanography, 2021, 66, 965-994. | 3.1 | 99 |
| 16 | Benthic protists: the under-charted majority. FEMS Microbiology Ecology, 2016, 92, fiw120. | 2.7 | 94 |
| 17 | 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 |
| 18 | Multigene phylogenetic analyses support recognition of the Sporolithales ord. nov Molecular Phylogenetics and Evolution, 2010, 54, 302-305. | 2.7 | 77 |

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|----|---|------|-----------|
| 19 | Communityâ€Level Responses to Iron Availability in Open Ocean Plankton Ecosystems. Global Biogeochemical Cycles, 2019, 33, 391-419. | 4.9 | 76 |
| 20 | Diversity patterns of uncultured Haptophytes unravelled by pyrosequencing in Naples Bay. Molecular Ecology, 2013, 22, 87-101. | 3.9 | 70 |
| 21 | 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 |
| 22 | Ocean acidification shows negligible impacts on high-latitude bacterial community structure in coastal pelagic mesocosms. Biogeosciences, 2013, 10, 555-566. | 3.3 | 60 |
| 23 | Mixotrophic protists display contrasted biogeographies in the global ocean. ISME Journal, 2019, 13, 1072-1083. | 9.8 | 55 |
| 24 | 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 |
| 25 | Some considerations for analyzing biodiversity using integrative metagenomics and gene networks. Biology Direct, 2010, 5, 47. | 4.6 | 50 |
| 26 | 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 |
| 27 | Clade-specific diversification dynamics of marine diatoms since the Jurassic. Nature Ecology and Evolution, 2018, 2, 1715-1723. | 7.8 | 40 |
| 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 | 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 |
| 30 | Acclimation of a low iron adapted Ostreococcus strain to iron limitation through cell biomass lowering. Scientific Reports, 2017, 7, 327. | 3.3 | 25 |
| 31 | A de novo approach to disentangle partner identity and function in holobiont systems. Microbiome, 2018, 6, 105. | 11.1 | 19 |
| 32 | Towards omics-based predictions of planktonic functional composition from environmental data. Nature Communications, 2021, 12, 4361. | 12.8 | 16 |
| 33 | A resource-frugal probabilistic dictionary and applications in bioinformatics. Discrete Applied Mathematics, 2020, 274, 92-102. | 0.9 | 13 |
| 34 | 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 |
| 35 | Mare Incognitum: A Glimpse into Future Plankton Diversity and Ecology Research. Frontiers in Marine Science, 2017, 4, . | 2.5 | 10 |