Cynthia B Silveira

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6957000/publications.pdf

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36 papers 2,198 citations

304743 22 h-index 35 g-index

44 all docs

44 docs citations

times ranked

44

2756 citing authors

#	Article	IF	CITATIONS
1	Distribution of soil viruses across China and their potential role in phosphorous metabolism. Environmental Microbiomes, 2022, 17, 6.	5.0	17
2	Benthic assemblages are more predictable than fish assemblages at an island scale. Coral Reefs, 2022, 41, 1031-1043.	2.2	3
3	Nasopharyngeal Microbial Communities of Patients Infected With SARS-CoV-2 That Developed COVID-19. Frontiers in Microbiology, 2021, 12, 637430.	3. 5	53
4	Three-Dimensional Molecular Cartography of the Caribbean Reef-Building Coral Orbicella faveolata. Frontiers in Marine Science, 2021, 8, .	2.5	11
5	Prophage Genomics and Ecology in the Family Rhodobacteraceae. Microorganisms, 2021, 9, 1115.	3.6	22
6	The landscape of lysogeny across microbial community density, diversity and energetics. Environmental Microbiology, 2021, 23, 4098-4111.	3.8	50
7	Space-filling and benthic competition on coral reefs. PeerJ, 2021, 9, e11213.	2.0	7
8	Extending the natural adaptive capacity of coral holobionts. Nature Reviews Earth & Environment, 2021, 2, 747-762.	29.7	110
9	Multi-Omics Study of Keystone Species in a Cystic Fibrosis Microbiome. International Journal of Molecular Sciences, 2021, 22, 12050.	4.1	14
10	Quantification of Lysogeny Caused by Phage Coinfections in Microbial Communities from Biophysical Principles. MSystems, 2020, 5, .	3.8	28
11	A multiomic analysis of in situ coral–turf algal interactions. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 13588-13595.	7.1	48
12	Genomic and ecological attributes of marine bacteriophages encoding bacterial virulence genes. BMC Genomics, 2020, 21, 126.	2.8	26
13	Reply to: Caution in inferring viral strategies from abundance correlations in marine metagenomes. Nature Communications, 2019, 10, 502.	12.8	2
14	Biophysical and physiological processes causing oxygen loss from coral reefs. ELife, 2019, 8, .	6.0	19
15	Release of dissolved and particulate organic matter by the soft coral Lobophytum and subsequent microbial degradation. Journal of Experimental Marine Biology and Ecology, 2018, 504, 53-60.	1.5	10
16	Microbial processes driving coral reef organic carbon flow. FEMS Microbiology Reviews, 2017, 41, 575-595.	8.6	67
17	Variability and host density independence in inductions-based estimates of environmental lysogeny. Nature Microbiology, 2017, 2, 17064.	13.3	57
18	Marine viruses discovered via metagenomics shed light on viral strategies throughout the oceans. Nature Communications, 2017, 8, 15955.	12.8	231

#	Article	IF	Citations
19	Bacterial Community Associated with the Reef Coral Mussismilia braziliensis's Momentum Boundary Layer over a Diel Cycle. Frontiers in Microbiology, 2017, 8, 784.	3.5	30
20	Piggyback-the-Winner in host-associated microbial communities. Npj Biofilms and Microbiomes, 2016, 2, 16010.	6.4	189
21	Energetic differences between bacterioplankton trophic groups and coral reef resistance. Proceedings of the Royal Society B: Biological Sciences, 2016, 283, 20160467.	2.6	30
22	Global microbialization of coral reefs. Nature Microbiology, 2016, 1, 16042.	13.3	214
23	Lytic to temperate switching of viral communities. Nature, 2016, 531, 466-470.	27.8	440
24	Microbial and sponge loops modify fish production in phaseâ€shifting coral reefs. Environmental Microbiology, 2015, 17, 3832-3846.	3.8	43
25	Physiologic and metagenomic attributes of the rhodoliths forming the largest CaCO3 bed in the South Atlantic Ocean. ISME Journal, 2014, 8, 52-62.	9.8	68
26	The Family Alcanivoraceae., 2014,, 59-67.		0
27	Antibiotic Resistance is Widespread in Urban Aquatic Environments of Rio de Janeiro, Brazil. Microbial Ecology, 2014, 68, 441-452.	2.8	33
28	Microbial Community Compositional Shifts in Bleached Colonies of the Brazilian Reef-Building Coral Siderastrea stellata. Microbial Ecology, 2013, 65, 205-213.	2.8	22
29	Bacterial community composition shifts in the gut of Periplaneta americana fed on different lignocellulosic materials. SpringerPlus, 2013, 2, 609.	1.2	50
30	Tropical Aquatic Archaea Show Environment-Specific Community Composition. PLoS ONE, 2013, 8, e76321.	2.5	10
31	Influence of Salinity on Bacterioplankton Communities from the Brazilian Rain Forest to the Coastal Atlantic Ocean. PLoS ONE, 2011, 6, e17789.	2.5	36
32	Archaea, Bacteria, and Algal Plastids Associated with the Reef-Building Corals Siderastrea stellata and Mussismilia hispida from Búzios, South Atlantic Ocean, Brazil. Microbial Ecology, 2010, 59, 523-532.	2.8	40
33	Environmental Shaping of Sponge Associated Archaeal Communities. PLoS ONE, 2010, 5, e15774.	2.5	84
34	Archaeal and bacterial communities of heavy metal contaminated acidic waters from zinc mine residues in Sepetiba Bay. Extremophiles, 2009, 13, 263-271.	2.3	54
35	Bacterial communities of the marine sponges Hymeniacidon heliophila and Polymastia janeirensis and their environment in Rio de Janeiro, Brazil. Marine Biology, 2008, 155, 135-146.	1.5	25
36	Prokaryotic diversity in one of the largest hypersaline coastal lagoons in the world. Extremophiles, 2008, 12, 595-604.	2.3	34