Sara Beier

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8956007/publications.pdf

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567144 610775 1,186 25 15 24 citations h-index g-index papers 1925 27 27 27 citing authors all docs docs citations times ranked

#	Article	IF	CITATIONS
1	Bacterial chitin degradation—mechanisms and ecophysiological strategies. Frontiers in Microbiology, 2013, 4, 149.	1.5	368
2	Function-specific response to depletion of microbial diversity. ISME Journal, 2011, 5, 351-361.	4.4	183
3	Bacterial Community Composition in Central European Running Waters Examined by Temperature Gradient Gel Electrophoresis and Sequence Analysis of 16S rRNA Genes. Applied and Environmental Microbiology, 2008, 74, 188-199.	1.4	69
4	Ecosystem-wide metagenomic binning enables prediction of ecological niches from genomes. Communications Biology, 2020, 3, 119.	2.0	64
5	Uncoupling of chitinase activity and uptake of hydrolysis products in freshwater bacterioplankton. Limnology and Oceanography, 2011, 56, 1179-1188.	1.6	62
6	Effect of large magnetotactic bacteria with polyphosphate inclusions on the phosphate profile of the suboxic zone in the Black Sea. ISME Journal, 2019, 13, 1198-1208.	4.4	59
7	The transcriptional response of prokaryotes to phytoplanktonâ€derived dissolved organic matter in seawater. Environmental Microbiology, 2015, 17, 3466-3480.	1.8	55
8	BARM and BalticMicrobeDB, a reference metagenome and interface to meta-omic data for the Baltic Sea. Scientific Data, 2018, 5, 180146.	2.4	54
9	Interactions of Freshwater Cyanobacteria with Bacterial Antagonists. Applied and Environmental Microbiology, 2017, 83, .	1.4	39
10	Experimental insights into the importance of ecologically dissimilar bacteria to community assembly along a salinity gradient. Environmental Microbiology, 2018, 20, 1170-1184.	1.8	32
11	Microbial iron metabolism as revealed by gene expression profiles in contrasted Southern Ocean regimes. Environmental Microbiology, 2019, 21, 2360-2374.	1.8	27
12	High Ratio of Bacteriochlorophyll Biosynthesis Genes to Chlorophyll Biosynthesis Genes in Bacteria of Humic Lakes. Applied and Environmental Microbiology, 2009, 75, 7221-7228.	1.4	23
13	Global Phylogeography of Chitinase Genes in Aquatic Metagenomes. Applied and Environmental Microbiology, 2011, 77, 1101-1106.	1.4	21
14	The transcriptional regulation of the glyoxylate cycle in <scp>SAR</scp> 11 in response to iron fertilization in the <scp>S</scp> outhern <scp>O</scp> cean. Environmental Microbiology Reports, 2015, 7, 427-434.	1.0	20
15	Phenotypic plasticity in heterotrophic marine microbial communities in continuous cultures. ISME Journal, 2015, 9, 1141-1151.	4.4	20
16	Metatranscriptomic data reveal the effect of different community properties on multifunctional redundancy. Molecular Ecology, 2017, 26, 6813-6826.	2.0	18
17	Pronounced seasonal dynamics of freshwater chitinase genes and chitin processing. Environmental Microbiology, 2012, 14, 2467-2479.	1.8	12
18	Uptake of Leucine, Chitin, and Iron by Prokaryotic Groups during Spring Phytoplankton Blooms Induced by Natural Iron Fertilization off Kerguelen Island (Southern Ocean). Frontiers in Marine Science, 2016, 3, .	1.2	12

#	Article	IF	CITATIONS
19	Lake bacterioplankton dynamics over diurnal timescales. Freshwater Biology, 2017, 62, 191-204.	1.2	11
20	A metatranscriptomicsâ€based assessment of smallâ€scale mixing of sulfidic and oxic waters on redoxcline prokaryotic communities. Environmental Microbiology, 2018, 21, 584-602.	1.8	10
21	Niche breadth affects bacterial transcription patterns along a salinity gradient. Molecular Ecology, 2022, 31, 1216-1233.	2.0	10
22	Cryopreservation and Resuscitation of Natural Aquatic Prokaryotic Communities. Frontiers in Microbiology, 2020, 11, 597653.	1.5	8
23	The environment drives microbial trait variability in aquatic habitats. Molecular Ecology, 2020, 29, 4605-4617.	2.0	5
24	Betaproteobacterial ammonia oxidizers in root zones of aquatic macrophytes. Fundamental and Applied Limnology, 2010, 177, 241-255.	0.4	3
25	Editorial: Advancements in the Understanding of Anthropogenic Impacts on the Microbial Ecology and Function of Aquatic Environments. Frontiers in Microbiology, 2021, 12, 820697.	1.5	0