

Jennifer B H Martiny

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3551897/publications.pdf>

Version: 2024-02-01

93
papers

19,015
citations

57758

44
h-index

48315

88
g-index

101
all docs

101
docs citations

101
times ranked

20158
citing authors

#	ARTICLE	IF	CITATIONS
1	Differential Response of Bacterial Microdiversity to Simulated Global Change. <i>Applied and Environmental Microbiology</i> , 2022, 88, aem0242921.	3.1	7
2	Bacterial community response to environmental change varies with depth in the surface soil. <i>Soil Biology and Biochemistry</i> , 2022, 172, 108761.	8.8	15
3	Routes and rates of bacterial dispersal impact surface soil microbiome composition and functioning. <i>ISME Journal</i> , 2022, 16, 2295-2304.	9.8	26
4	Microbial community response to a decade of simulated global changes depends on the plant community. <i>Elementa</i> , 2021, 9, .	3.2	10
5	Adaptive differentiation and rapid evolution of a soil bacterium along a climate gradient. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	56
6	High-Fiber, Whole-Food Dietary Intervention Alters the Human Gut Microbiome but Not Fecal Short-Chain Fatty Acids. <i>MSystems</i> , 2021, 6, .	3.8	69
7	Defining trait-based microbial strategies with consequences for soil carbon cycling under climate change. <i>ISME Journal</i> , 2020, 14, 1-9.	9.8	470
8	Evolutionary relationships among bifidobacteria and their hosts and environments. <i>BMC Genomics</i> , 2020, 21, 26.	2.8	26
9	The emergence of microbiome centres. <i>Nature Microbiology</i> , 2020, 5, 2-3.	13.3	13
10	Alpha-, beta-, and gamma-diversity of bacteria varies across habitats. <i>PLoS ONE</i> , 2020, 15, e0233872.	2.5	105
11	Cervicovaginal Microbiome Composition Is Associated with Metabolic Profiles in Healthy Pregnancy. <i>MBio</i> , 2020, 11, .	4.1	30
12	Fiber Force: A Fiber Diet Intervention in an Advanced Course-Based Undergraduate Research Experience (CURE) Course. <i>Journal of Microbiology and Biology Education</i> , 2020, 21, .	1.0	15
13	Phylogenetic conservation of soil bacterial responses to simulated global changes. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2020, 375, 20190242.	4.0	46
14	Conceptual challenges in microbial community ecology. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2020, 375, 20190241.	4.0	21
15	Drought and plant litter chemistry alter microbial gene expression and metabolite production. <i>ISME Journal</i> , 2020, 14, 2236-2247.	9.8	79
16	Comparative Genomics of Nitrogen Cycling Pathways in Bacteria and Archaea. <i>Microbial Ecology</i> , 2019, 77, 597-606.	2.8	21
17	Scientists's warning to humanity: microorganisms and climate change. <i>Nature Reviews Microbiology</i> , 2019, 17, 569-586.	28.6	1,138
18	Phylogenetic conservation of bacterial responses to soil nitrogen addition across continents. <i>Nature Communications</i> , 2019, 10, 2499.	12.8	48

#	ARTICLE	IF	CITATIONS
19	Optimization of a Method To Quantify Soil Bacterial Abundance by Flow Cytometry. <i>MSphere</i> , 2019, 4, .	2.9	25
20	Maintenance of Sympatric and Allopatric Populations in Free-Living Terrestrial Bacteria. <i>MBio</i> , 2019, 10, .	4.1	19
21	Experimental Evidence that Stochasticity Contributes to Bacterial Composition and Functioning in a Decomposer Community. <i>MBio</i> , 2019, 10, .	4.1	23
22	Towards a Natural History of Soil Bacterial Communities. <i>Trends in Microbiology</i> , 2018, 26, 250-252.	7.7	7
23	Microbial decomposers not constrained by climate history along a Mediterranean climate gradient in southern California. <i>Ecology</i> , 2018, 99, 1441-1452.	3.2	16
24	Bacterial diversity is positively correlated with soil heterogeneity. <i>Ecosphere</i> , 2018, 9, e02079.	2.2	68
25	Dispersal alters bacterial diversity and composition in a natural community. <i>ISME Journal</i> , 2018, 12, 296-299.	9.8	70
26	Decomposition responses to climate depend on microbial community composition. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 11994-11999.	7.1	214
27	Emergence of soil bacterial ecotypes along a climate gradient. <i>Environmental Microbiology</i> , 2018, 20, 4112-4126.	3.8	32
28	The importance of resolving biogeographic patterns of microbial microdiversity. <i>Microbiology Australia</i> , 2018, 39, 5.	0.4	23
29	Is Throwing an Apple Core Out of the Car Littering?â€”Microbial Communities in Natural Composting. <i>Frontiers for Young Minds</i> , 2018, 6, .	0.8	0
30	Broadscale Ecological Patterns Are Robust to Use of Exact Sequence Variants versus Operational Taxonomic Units. <i>MSphere</i> , 2018, 3, .	2.9	168
31	Predictable Molecular Adaptation of Coevolving <i>Enterococcus faecium</i> and Lytic Phage EfV12-phi1. <i>Frontiers in Microbiology</i> , 2018, 9, 3192.	3.5	30
32	The effect of soil inoculants on seed germination of native and invasive species. <i>Botany</i> , 2017, 95, 469-480.	1.0	16
33	Microdiversity of an Abundant Terrestrial Bacterium Encompasses Extensive Variation in Ecologically Relevant Traits. <i>MBio</i> , 2017, 8, .	4.1	49
34	Microbial legacies alter decomposition in response to simulated global change. <i>ISME Journal</i> , 2017, 11, 490-499.	9.8	112
35	Effects of dispersal and selection on stochastic assembly in microbial communities. <i>ISME Journal</i> , 2017, 11, 176-185.	9.8	256
36	Phylogenetic conservation of substrate use specialization in leaf litter bacteria. <i>PLoS ONE</i> , 2017, 12, e0174472.	2.5	14

#	ARTICLE	IF	CITATIONS
37	Biogeographic Variation in Host Range Phenotypes and Taxonomic Composition of Marine Cyanophage Isolates. <i>Frontiers in Microbiology</i> , 2016, 7, 983.	3.5	26
38	Evidence for Ecological Flexibility in the Cosmopolitan Genus <i>Curtobacterium</i> . <i>Frontiers in Microbiology</i> , 2016, 7, 1874.	3.5	66
39	Genomic diversification of marine cyanophages into stable ecotypes. <i>Environmental Microbiology</i> , 2016, 18, 4240-4253.	3.8	44
40	Global biogeography of microbial nitrogen-cycling traits in soil. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 8033-8040.	7.1	365
41	The genomic content and context of auxiliary metabolic genes in marine cyanomyoviruses. <i>Virology</i> , 2016, 499, 219-229.	2.4	99
42	The Microbial Olympics 2016. <i>Nature Microbiology</i> , 2016, 1, 16122.	13.3	7
43	History Leaves Its Mark on Soil Bacterial Diversity. <i>MBio</i> , 2016, 7, .	4.1	14
44	Microbial response to simulated global change is phylogenetically conserved and linked with functional potential. <i>ISME Journal</i> , 2016, 10, 109-118.	9.8	123
45	Nitrification kinetics and ammonia-oxidizing community respond to warming and altered precipitation. <i>Ecosphere</i> , 2015, 6, 1-17.	2.2	19
46	Nonlinear responses in salt marsh functioning to increased nitrogen addition. <i>Ecology</i> , 2015, 96, 936-947.	3.2	31
47	Temporal variation overshadows the response of leaf litter microbial communities to simulated global change. <i>ISME Journal</i> , 2015, 9, 2477-2489.	9.8	112
48	Nitrogen addition, not initial phylogenetic diversity, increases litter decomposition by fungal communities. <i>Frontiers in Microbiology</i> , 2015, 6, 109.	3.5	17
49	Microbiomes in light of traits: A phylogenetic perspective. <i>Science</i> , 2015, 350, aac9323.	12.6	652
50	Nitrogen Cycling Potential of a Grassland Litter Microbial Community. <i>Applied and Environmental Microbiology</i> , 2015, 81, 7012-7022.	3.1	51
51	Microbial composition alters the response of litter decomposition to environmental change. <i>Ecology</i> , 2015, 96, 154-163.	3.2	71
52	Dispersal and the Microbiome. <i>Microbe Magazine</i> , 2015, 10, 191-196.	0.4	13
53	Cellulolytic potential under environmental changes in microbial communities from grassland litter. <i>Frontiers in Microbiology</i> , 2014, 5, 639.	3.5	61
54	Antagonistic Coevolution of Marine Planktonic Viruses and Their Hosts. <i>Annual Review of Marine Science</i> , 2014, 6, 393-414.	11.6	68

#	ARTICLE	IF	CITATIONS
55	Abundance of Broad Bacterial Taxa in the Sargasso Sea Explained by Environmental Conditions but Not Water Mass. <i>Applied and Environmental Microbiology</i> , 2014, 80, 2786-2795.	3.1	36
56	Relationships between Methylobacteria and Glyphosate with Native and Invasive Plant Species: Implications for Restoration. <i>Restoration Ecology</i> , 2013, 21, 105-113.	2.9	12
57	Beta diversity of marine bacteria depends on temporal scale. <i>Ecology</i> , 2013, 94, 1898-1904.	3.2	75
58	Microbial abundance and composition influence litter decomposition response to environmental change. <i>Ecology</i> , 2013, 94, 714-725.	3.2	340
59	Marine cyanophages exhibit local and regional biogeography. <i>Environmental Microbiology</i> , 2013, 15, 1452-1463.	3.8	43
60	Microbial composition affects the functioning of estuarine sediments. <i>ISME Journal</i> , 2013, 7, 868-879.	9.8	130
61	Macroecological patterns of marine bacteria on a global scale. <i>Journal of Biogeography</i> , 2013, 40, 800-811.	3.0	53
62	Microbial Biodiversity. , 2013, , 252-258.		2
63	Microbial Biogeography. , 2013, , 271-279.		1
64	Coupled high-throughput functional screening and next generation sequencing for identification of plant polymer decomposing enzymes in metagenomic libraries. <i>Frontiers in Microbiology</i> , 2013, 4, 282.	3.5	44
65	Rapid diversification of coevolving marine <i>Synechococcus</i> and a virus. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 4544-4549.	7.1	178
66	Fundamentals of Microbial Community Resistance and Resilience. <i>Frontiers in Microbiology</i> , 2012, 3, 417.	3.5	1,131
67	An atomic force microscopy investigation of cyanophage structure. <i>Micron</i> , 2012, 43, 1336-1342.	2.2	9
68	The Abundance of Pink-Pigmented Facultative Methylophages in the Root Zone of Plant Species in Invaded Coastal Sage Scrub Habitat. <i>PLoS ONE</i> , 2012, 7, e31026.	2.5	15
69	The Effect of Nitrogen Enrichment on C1-Cycling Microorganisms and Methane Flux in Salt Marsh Sediments. <i>Frontiers in Microbiology</i> , 2012, 3, 90.	3.5	24
70	Beyond biogeographic patterns: processes shaping the microbial landscape. <i>Nature Reviews Microbiology</i> , 2012, 10, 497-506.	28.6	1,299
71	Drivers of bacterial β -diversity depend on spatial scale. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 7850-7854.	7.1	672
72	Functional Metagenomics Reveals Previously Unrecognized Diversity of Antibiotic Resistance Genes in Gulls. <i>Frontiers in Microbiology</i> , 2011, 2, 238.	3.5	46

#	ARTICLE	IF	CITATIONS
73	Global Patterns of Bacterial Beta-Diversity in Seafloor and Seawater Ecosystems. PLoS ONE, 2011, 6, e24570.	2.5	525
74	Patterns of fungal diversity and composition along a salinity gradient. ISME Journal, 2011, 5, 379-388.	9.8	160
75	Nitrogen and phosphorus enrichment alter the composition of ammonia-oxidizing bacteria in salt marsh sediments. ISME Journal, 2010, 4, 933-944.	9.8	41
76	Structural analysis of a Synechococcus myovirus S-CAM4 and infected cells by atomic force microscopy. Journal of General Virology, 2010, 91, 3095-3104.	2.9	10
77	The minimum information about a genome sequence (MIGS) specification. Nature Biotechnology, 2008, 26, 541-547.	17.5	1,069
78	Pathogens promote plant diversity through a compensatory response. Ecology Letters, 2008, 11, 461-469.	6.4	71
79	Rapid evolution buffers ecosystem impacts of viruses in a microbial food web. Ecology Letters, 2008, 11, 1178-1188.	6.4	73
80	It's all relative: ranking the diversity of aquatic bacterial communities. Environmental Microbiology, 2008, 10, 2200-2210.	3.8	159
81	Resistance, resilience, and redundancy in microbial communities. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 11512-11519.	7.1	2,195
82	Selection and Characterization of Cyanophage Resistance in Marine <i>Synechococcus</i> Strains. Applied and Environmental Microbiology, 2007, 73, 5516-5522.	3.1	64
83	A COMPARISON OF TAXON CO-OCCURRENCE PATTERNS FOR MACRO- AND MICROORGANISMS. Ecology, 2007, 88, 1345-1353.	3.2	223
84	Is there a cost of virus resistance in marine cyanobacteria?. ISME Journal, 2007, 1, 300-312.	9.8	127
85	Testing the functional significance of microbial composition in natural communities. FEMS Microbiology Ecology, 2007, 62, 161-170.	2.7	173
86	Microbial biodiversity. , 2007, , 1-9.		1
87	Alkenone producers inferred from well-preserved 18S rDNA in Greenland lake sediments. Journal of Geophysical Research, 2006, 111, .	3.3	39
88	Microbial biogeography: putting microorganisms on the map. Nature Reviews Microbiology, 2006, 4, 102-112.	28.6	2,434
89	A taxonomic area relationship for bacteria. Nature, 2004, 432, 750-753.	27.8	632
90	Conservation of tropical forest birds in countryside habitats. Ecology Letters, 2002, 5, 121-129.	6.4	181

#	ARTICLE	IF	CITATIONS
91	Population Diversity, Overview. , 2001, , 168-174.		0
92	Counting the Uncountable: Statistical Approaches to Estimating Microbial Diversity. Applied and Environmental Microbiology, 2001, 67, 4399-4406.	3.1	1,032
93	Population Diversity: Its Extent and Extinction. Science, 1997, 278, 689-692.	12.6	471