

Eva S Lindström

List of Publications by Year in descending order

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Version: 2024-02-01

65
papers

6,028
citations

94415

37
h-index

110368

64
g-index

72
all docs

72
docs citations

72
times ranked

7001
citing authors

#	ARTICLE	IF	CITATIONS
1	Warming mediates the resistance of aquatic bacteria to invasion during community coalescence. <i>Molecular Ecology</i> , 2021, 30, 1345-1356.	3.9	9
2	High Iron Requirements for Growth in the Nuisance Alga <i>Gonyostomum semen</i> (Raphidophyceae). <i>Journal of Phycology</i> , 2021, 57, 1309-1322.	2.3	6
3	Comprehensive analysis of chemical and biological problems associated with browning agents used in aquatic studies. <i>Limnology and Oceanography: Methods</i> , 2021, 19, 818-835.	2.0	11
4	Streamlined and Abundant Bacterioplankton Thrive in Functional Cohorts. <i>MSystems</i> , 2020, 5, .	3.8	8
5	Different Roles of Environmental Selection, Dispersal, and Drift in the Assembly of Intestinal Microbial Communities of Freshwater Fish With and Without a Stomach. <i>Frontiers in Ecology and Evolution</i> , 2020, 8, .	2.2	10
6	Using null models to compare bacterial and microeukaryotic metacommunity assembly under shifting environmental conditions. <i>Scientific Reports</i> , 2020, 10, 2455.	3.3	64
7	Factors influencing aquatic and terrestrial bacterial community assembly. <i>Environmental Microbiology Reports</i> , 2019, 11, 306-315.	2.4	152
8	Sharing of photobionts in sympatric populations of <i>Thamnolia</i> and <i>Cetraria</i> lichens: evidence from high-throughput sequencing. <i>Scientific Reports</i> , 2018, 8, 4406.	3.3	29
9	Dispersal timing determines the importance of priority effects in bacterial communities. <i>ISME Journal</i> , 2018, 12, 644-646.	9.8	44
10	Decomposing multiple dimensions of stability in global change experiments. <i>Ecology Letters</i> , 2018, 21, 21-30.	6.4	167
11	High abundances of the nuisance raphidophyte <i>Gonyostomum semen</i> in brown water lakes are associated with high concentrations of iron. <i>Scientific Reports</i> , 2018, 8, 13463.	3.3	18
12	Repeated disturbances affect functional but not compositional resistance and resilience in an aquatic bacterioplankton community. <i>Environmental Microbiology Reports</i> , 2018, 10, 493-500.	2.4	18
13	Increased water colour affects freshwater plankton communities in a mesocosm study. <i>Aquatic Microbial Ecology</i> , 2018, 81, 1-17.	1.8	27
14	Effects of sterilization on dissolved organic carbon (DOC) composition and bacterial utilization of DOC from lakes. <i>Aquatic Microbial Ecology</i> , 2018, 82, 199-208.	1.8	8
15	Contribution of different bacterial dispersal sources to lakes: Population and community effects in different seasons. <i>Environmental Microbiology</i> , 2017, 19, 2391-2404.	3.8	50
16	Influence of pulsed and continuous substrate inputs on freshwater bacterial community composition and functioning in bioreactors. <i>Environmental Microbiology</i> , 2017, 19, 5078-5087.	3.8	7
17	Contribution of different dispersal sources to the metabolic response of lake bacterioplankton following a salinity change. <i>Environmental Microbiology</i> , 2017, 19, 251-260.	3.8	19
18	OVERVIEW Progress and perspectives in aquatic microbial ecology: highlights of the SAME 14, Uppsala, Sweden, 2015. <i>Aquatic Microbial Ecology</i> , 2017, 80, 101-103.	1.8	0

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19	Microbes as Engines of Ecosystem Function: When Does Community Structure Enhance Predictions of Ecosystem Processes?. <i>Frontiers in Microbiology</i> , 2016, 7, 214.	3.5	479
20	Remnants of marine bacterial communities can be retrieved from deep sediments in lakes of marine origin. <i>Environmental Microbiology Reports</i> , 2016, 8, 479-485.	2.4	10
21	Experimental insights into the importance of aquatic bacterial community composition to the degradation of dissolved organic matter. <i>ISME Journal</i> , 2016, 10, 533-545.	9.8	418
22	Combined effects of zooplankton grazing and dispersal on the diversity and assembly mechanisms of bacterial metacommunities. <i>Environmental Microbiology</i> , 2015, 17, 2275-2287.	3.8	47
23	Relationships between Bacterial Community Composition, Functional Trait Composition and Functioning Are Context Dependent “ but What Is the Context?. <i>PLoS ONE</i> , 2014, 9, e112409.	2.5	2
24	The spatial structure of bacterial communities is influenced by historical environmental conditions. <i>Ecology</i> , 2014, 95, 1134-1140.	3.2	67
25	Can marine bacteria be recruited from freshwater sources and the air?. <i>ISME Journal</i> , 2014, 8, 2423-2430.	9.8	55
26	Biogeography of bacterial communities exposed to progressive long-term environmental change. <i>ISME Journal</i> , 2013, 7, 937-948.	9.8	330
27	Unveiling Distribution Patterns of Freshwater Phytoplankton by a Next Generation Sequencing Based Approach. <i>PLoS ONE</i> , 2013, 8, e53516.	2.5	120
28	Variable Effects of Dispersal on Productivity of Bacterial Communities Due to Changes in Functional Trait Composition. <i>PLoS ONE</i> , 2013, 8, e80825.	2.5	20
29	Importance of space and the local environment for linking local and regional abundances of microbes. <i>Aquatic Microbial Ecology</i> , 2012, 67, 35-45.	1.8	8
30	Unraveling assembly of stream biofilm communities. <i>ISME Journal</i> , 2012, 6, 1459-1468.	9.8	242
31	Freshwater bacterioplankton richness in oligotrophic lakes depends on nutrient availability rather than on species-area relationships. <i>ISME Journal</i> , 2012, 6, 1127-1136.	9.8	105
32	Local and regional factors influencing bacterial community assembly. <i>Environmental Microbiology Reports</i> , 2012, 4, 1-9.	2.4	434
33	Which sequencing depth is sufficient to describe patterns in bacterial α - and β -diversity?. <i>Environmental Microbiology Reports</i> , 2012, 4, 367-372.	2.4	117
34	Function-specific response to depletion of microbial diversity. <i>ISME Journal</i> , 2011, 5, 351-361.	9.8	183
35	The Importance of Dispersal for Bacterial Community Composition and Functioning. <i>PLoS ONE</i> , 2011, 6, e25883.	2.5	82
36	Changing phosphorus concentration and subsequent prophage induction alter composition of a freshwater viral assemblage. <i>Freshwater Biology</i> , 2010, 55, 1984-1996.	2.4	6

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37	Species sorting affects bacterioplankton community composition as determined by 16S rDNA and 16S rRNA fingerprints. <i>ISME Journal</i> , 2010, 4, 729-738.	9.8	93
38	Regional invariance among microbial communities. <i>Ecology Letters</i> , 2010, 13, 118-127.	6.4	129
39	Ubiquity of <i>Polynucleobacter necessarius</i> ssp. <i>asymbioticus</i> in lentic freshwater habitats of a heterogeneous 2000 km ² area. <i>Environmental Microbiology</i> , 2010, 12, 658-669.	3.8	115
40	The interplay between bacterial community composition and the environment determining function of inland water bacteria. <i>Limnology and Oceanography</i> , 2010, 55, 2052-2060.	3.1	35
41	Temporal variation in freshwater viral and bacterial community composition. <i>Freshwater Biology</i> , 2008, 53, 1163-1175.	2.4	41
42	Variable importance of viral-induced bacterial mortality along gradients of trophic status and humic content in lakes. <i>Freshwater Biology</i> , 2008, 53, 1101-1113.	2.4	36
43	Composition and dispersal of riverine and lake phytoplankton communities in connected systems with different water retention times. <i>Freshwater Biology</i> , 2008, 53, 2520-2529.	2.4	8
44	Biogeography of Bacterioplankton in Inland Waters. <i>Freshwater Reviews: A Journal of the Freshwater Biological Association</i> , 2008, 1, 99-114.	1.0	106
45	DOES ECOSYSTEM SIZE DETERMINE AQUATIC BACTERIAL RICHNESS? COMMENT. <i>Ecology</i> , 2007, 88, 252-253.	3.2	16
46	First evidence for a bipolar distribution of dominant freshwater lake bacterioplankton. <i>Antarctic Science</i> , 2007, 19, 245-252.	0.9	38
47	THE ROLE OF ENVIRONMENTAL AND SPATIAL PROCESSES IN STRUCTURING LAKE COMMUNITIES FROM BACTERIA TO FISH. <i>Ecology</i> , 2006, 87, 2985-2991.	3.2	446
48	External control of bacterial community structure in lakes. <i>Limnology and Oceanography</i> , 2006, 51, 339-342.	3.1	108
49	Influence of dissolved organic matter source on lake bacterioplankton structure and function – implications for seasonal dynamics of community composition. <i>FEMS Microbiology Ecology</i> , 2006, 56, 406-417.	2.7	115
50	Structure and Function of Bacterial Communities Emerging from Different Sources under Identical Conditions. <i>Applied and Environmental Microbiology</i> , 2006, 72, 212-220.	3.1	155
51	Community composition of bacterioplankton and cell transport in lakes in two different drainage areas. <i>Aquatic Sciences</i> , 2005, 67, 210-219.	1.5	42
52	Weak coupling between community composition and functioning of aquatic bacteria. <i>Limnology and Oceanography</i> , 2005, 50, 957-967.	3.1	170
53	Distribution of Typical Freshwater Bacterial Groups Is Associated with pH, Temperature, and Lake Water Retention Time. <i>Applied and Environmental Microbiology</i> , 2005, 71, 8201-8206.	3.1	402
54	Production and food web interactions of Arctic freshwater plankton and responses to increased DOC. <i>Archiv für Hydrobiologie</i> , 2004, 159, 289-307.	1.1	19

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55	Response of a member of the Verrucomicrobia, among the dominating bacteria in a hypolimnion, to increased phosphorus availability. <i>Journal of Plankton Research</i> , 2004, 26, 241-246.	1.8	56
56	Influence of inlet bacteria on bacterioplankton assemblage composition in lakes of different hydraulic retention time. <i>Limnology and Oceanography</i> , 2004, 49, 125-136.	3.1	87
57	Growth dynamics within bacterial communities in riverine and estuarine batch cultures. <i>Aquatic Microbial Ecology</i> , 2004, 37, 137-148.	1.8	16
58	Viral and Bacterioplankton Dynamics in Two Lakes with Different Humic Contents. <i>Microbial Ecology</i> , 2003, 46, 406-415.	2.8	40
59	Rapid Screening for Freshwater Bacterial Groups by Using Reverse Line Blot Hybridization. <i>Applied and Environmental Microbiology</i> , 2003, 69, 5875-5883.	3.1	100
60	Live sorting and survival of unstained and DAPI-stained ciliates by flow cytometry. <i>Archiv für Hydrobiologie</i> , 2003, 157, 173-184.	1.1	1
61	Interactive effect of temperature and food concentration on growth rate: A test case using the small freshwater ciliate <i>Urotricha farcta</i> . <i>Limnology and Oceanography</i> , 2002, 47, 1447-1455.	3.1	69
62	Enumeration of small ciliates in culture by flow cytometry and nucleic acid staining. <i>Journal of Microbiological Methods</i> , 2002, 49, 173-182.	1.6	30
63	Do neighboring lakes share common taxa of bacterioplankton? Comparison of 16S rDNA fingerprints and sequences from three geographic regions. <i>Microbial Ecology</i> , 2002, 44, 1-9.	2.8	56
64	Investigating Influential Factors on Bacterioplankton Community Composition: Results from a Field Study of Five Mesotrophic Lakes. <i>Microbial Ecology</i> , 2001, 42, 598-605.	2.8	83
65	Bacterioplankton community composition in a boreal forest lake. <i>FEMS Microbiology Ecology</i> , 1998, 27, 163-174.	2.7	41