

James C Stegen

List of Publications by Year in descending order

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Version: 2024-02-01

115
papers

14,596
citations

61857

43
h-index

22764

112
g-index

153
all docs

153
docs citations

153
times ranked

16271
citing authors

#	ARTICLE	IF	CITATIONS
1	Distinct and Temporally Stable Assembly Mechanisms Shape Bacterial and Fungal Communities in Vineyard Soils. <i>Microbial Ecology</i> , 2023, 86, 337-349.	1.4	6
2	ORT: a workflow linking genome-scale metabolic models with reactive transport codes. <i>Bioinformatics</i> , 2022, 38, 778-784.	1.8	2
3	The ecological assembly of bacterial communities in Antarctic wetlands varies across levels of phylogenetic resolution. <i>Environmental Microbiology</i> , 2022, , .	1.8	1
4	Implications of sample treatment on characterization of riverine dissolved organic matter. <i>Environmental Sciences: Processes and Impacts</i> , 2022, 24, 773-782.	1.7	6
5	Inferring the Contribution of Microbial Taxa and Organic Matter Molecular Formulas to Ecological Assembly. <i>Frontiers in Microbiology</i> , 2022, 13, 803420.	1.5	5
6	Integrated, Coordinated, Open, and Networked (ICON) Science to Advance the Geosciences: Introduction and Synthesis of a Special Collection of Commentary Articles. <i>Earth and Space Science</i> , 2022, 9, .	1.1	14
7	Movement with meaning: integrating information into meta-ecology. <i>Oikos</i> , 2022, 2022, .	1.2	12
8	Advancing river corridor science beyond disciplinary boundaries with an inductive approach to catalyse hypothesis generation. <i>Hydrological Processes</i> , 2022, 36, .	1.1	7
9	Dissolved oxygen sensor in an automated hyporheic sampling system reveals biogeochemical dynamics. , 2022, 1, e0000014.		0
10	Disinfection byproducts formed during drinking water treatment reveal an export control point for dissolved organic matter in a subalpine headwater stream. <i>Water Research X</i> , 2022, 15, 100144.	2.8	7
11	Hot Spots and Hot Moments in the Critical Zone: Identification of and Incorporation into Reactive Transport Models. , 2022, , 9-47.		7
12	It Takes a Village: Using a Crowdsourced Approach to Investigate Organic Matter Composition in Global Rivers Through the Lens of Ecological Theory. <i>Frontiers in Water</i> , 2022, 4, .	1.0	3
13	Riverbed Temperature and 4D ERT Monitoring Reveals Heterogenous Horizontal and Vertical Groundwater-Surface Water Exchange Flows Under Dynamic Stage Conditions. <i>Frontiers in Earth Science</i> , 2022, 10, .	0.8	1
14	Continental-scale niche differentiation of dominant topsoil archaea in drylands. <i>Environmental Microbiology</i> , 2022, 24, 5483-5497.	1.8	3
15	Microbial and Environmental Processes Shape the Link between Organic Matter Functional Traits and Composition. <i>Environmental Science & Technology</i> , 2022, 56, 10504-10516.	4.6	27
16	Organic matter transformations are disconnected between surface water and the hyporheic zone. <i>Biogeosciences</i> , 2022, 19, 3099-3110.	1.3	4
17	Integrating field observations and process-based modeling to predict watershed water quality under environmental perturbations. <i>Journal of Hydrology</i> , 2021, 602, 125762.	2.3	22
18	A genomic catalog of Earth's microbiomes. <i>Nature Biotechnology</i> , 2021, 39, 499-509.	9.4	457

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19	Small streams dominate US tidal reaches and will be disproportionately impacted by sea-level rise. <i>Science of the Total Environment</i> , 2021, 753, 141944.	3.9	7
20	Special Collection on Open Collaboration Across Geosciences. <i>Eos</i> , 2021, 102, .	0.1	20
21	Coupled Biotic-Abiotic Processes Control Biogeochemical Cycling of Dissolved Organic Matter in the Columbia River Hyporheic Zone. <i>Frontiers in Water</i> , 2021, 2, .	1.0	18
22	Antecedent conditions determine the biogeochemical response of coastal soils to seawater exposure. <i>Soil Biology and Biochemistry</i> , 2021, 153, 108104.	4.2	7
23	Historical Contingency in Microbial Resilience to Hydrologic Perturbations. <i>Frontiers in Water</i> , 2021, 3, .	1.0	2
24	Evaluating a Laboratory Flume Microbiome as a Window Into Natural Riverbed Biogeochemistry. <i>Frontiers in Water</i> , 2021, 3, .	1.0	3
25	Sample Identifiers and Metadata to Support Data Management and Reuse in Multidisciplinary Ecosystem Sciences. <i>Data Science Journal</i> , 2021, 20, 11.	0.6	11
26	Assembly of the <i>Populus</i> Microbiome Is Temporally Dynamic and Determined by Selective and Stochastic Factors. <i>MSphere</i> , 2021, 6, e0131620.	1.3	25
27	Disturbance triggers non-linear microbe–environment feedbacks. <i>Biogeosciences</i> , 2021, 18, 4773-4789.	1.3	8
28	Amount and reactivity of dissolved organic matter export are affected by land cover change from old-growth to second-growth forests in headwater ecosystems. <i>Hydrological Processes</i> , 2021, 35, e14343.	1.1	3
29	A novel construct for scaling groundwater–river interactions based on machine-guided hydromorphic classification. <i>Environmental Research Letters</i> , 2021, 16, 104016.	2.2	1
30	Ecological theory applied to environmental metabolomes reveals compositional divergence despite conserved molecular properties. <i>Science of the Total Environment</i> , 2021, 788, 147409.	3.9	21
31	Contrasting Community Assembly Forces Drive Microbial Structural and Potential Functional Responses to Precipitation in an Incipient Soil System. <i>Frontiers in Microbiology</i> , 2021, 12, 754698.	1.5	4
32	TRY plant trait database – enhanced coverage and open access. <i>Global Change Biology</i> , 2020, 26, 119-188.	4.2	1,038
33	Temperature drives local contributions to beta diversity in mountain streams: Stochastic and deterministic processes. <i>Global Ecology and Biogeography</i> , 2020, 29, 420-432.	2.7	30
34	Tree growth, transpiration, and water-use efficiency between shoreline and upland red maple (<i>Acer</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5	1.9	10
35	Representing Organic Matter Thermodynamics in Biogeochemical Reactions via Substrate-Explicit Modeling. <i>Frontiers in Microbiology</i> , 2020, 11, 531756.	1.5	27
36	Using Community Science to Reveal the Global Chemogeography of River Metabolomes. <i>Metabolites</i> , 2020, 10, 518.	1.3	27

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37	Using metacommunity ecology to understand environmental metabolomes. <i>Nature Communications</i> , 2020, 11, 6369.	5.8	51
38	Carbon Limitation Leads to Thermodynamic Regulation of Aerobic Metabolism. <i>Environmental Science and Technology Letters</i> , 2020, 7, 517-524.	3.9	32
39	Straw chemistry links the assembly of bacterial communities to decomposition in paddy soils. <i>Soil Biology and Biochemistry</i> , 2020, 148, 107866.	4.2	49
40	Distinct assembly mechanisms underlie similar biogeographical patterns of rare and abundant bacteria in Tibetan Plateau grassland soils. <i>Environmental Microbiology</i> , 2020, 22, 2261-2272.	1.8	77
41	Localized basal area affects soil respiration temperature sensitivity in a coastal deciduous forest. <i>Biogeosciences</i> , 2020, 17, 771-780.	1.3	5
42	Ecological Assembly Processes Are Coordinated between Bacterial and Viral Communities in Fractured Shale Ecosystems. <i>MSystems</i> , 2020, 5, .	1.7	15
43	Distinct temporal diversity profiles for nitrogen cycling genes in a hyporheic microbiome. <i>PLoS ONE</i> , 2020, 15, e0228165.	1.1	12
44	Methane and nitrous oxide porewater concentrations and surface fluxes of a regulated river. <i>Science of the Total Environment</i> , 2020, 715, 136920.	3.9	20
45	A Flux Detection Probe to Quantify Dynamic Groundwater-Surface Water Exchange in the Hyporheic Zone. <i>Ground Water</i> , 2020, 58, 892-900.	0.7	8
46	Active layer depth and soil properties impact specific leaf area variation and ecosystem productivity in a boreal forest. <i>PLoS ONE</i> , 2020, 15, e0232506.	1.1	8
47	Spatial gradients in the characteristics of soil-carbon fractions are associated with abiotic features but not microbial communities. <i>Biogeosciences</i> , 2019, 16, 3911-3928.	1.3	19
48	Forfeiting the priority effect: turnover defines biofilm community succession. <i>ISME Journal</i> , 2019, 13, 1865-1877.	4.4	83
49	Assessing Microbial Community Patterns During Incipient Soil Formation From Basalt. <i>Journal of Geophysical Research G: Biogeosciences</i> , 2019, 124, 941-958.	1.3	16
50	Spatial and temporal variation in river corridor exchange across a 5th-order mountain stream network. <i>Hydrology and Earth System Sciences</i> , 2019, 23, 5199-5225.	1.9	23
51	Subsurface biogeochemistry is a missing link between ecology and hydrology in dam-impacted river corridors. <i>Science of the Total Environment</i> , 2019, 657, 435-445.	3.9	19
52	Co-located contemporaneous mapping of morphological, hydrological, chemical, and biological conditions in a 5th-order mountain stream network, Oregon, USA. <i>Earth System Science Data</i> , 2019, 11, 1567-1581.	3.7	14
53	Soil pH mediates the balance between stochastic and deterministic assembly of bacteria. <i>ISME Journal</i> , 2018, 12, 1072-1083.	4.4	591
54	Riverbed Hydrologic Exchange Dynamics in a Large Regulated River Reach. <i>Water Resources Research</i> , 2018, 54, 2715-2730.	1.7	17

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55	Influences of organic carbon speciation on hyporheic corridor biogeochemistry and microbial ecology. <i>Nature Communications</i> , 2018, 9, 585.	5.8	110
56	WHONDORS: a Community Resource for Studying Dynamic River Corridors. <i>MSystems</i> , 2018, 3, .	1.7	22
57	Drought Conditions Maximize the Impact of High-Frequency Flow Variations on Thermal Regimes and Biogeochemical Function in the Hyporheic Zone. <i>Water Resources Research</i> , 2018, 54, 7361-7382.	1.7	63
58	Two key features influencing community assembly processes at regional scale: Initial state and degree of change in environmental conditions. <i>Molecular Ecology</i> , 2018, 27, 5238-5251.	2.0	147
59	At the Nexus of History, Ecology, and Hydrobiogeochemistry: Improved Predictions across Scales through Integration. <i>MSystems</i> , 2018, 3, .	1.7	5
60	A unified conceptual framework for prediction and control of microbiomes. <i>Current Opinion in Microbiology</i> , 2018, 44, 20-27.	2.3	42
61	Multi 'omics comparison reveals metabolome biochemistry, not microbiome composition or gene expression, corresponds to elevated biogeochemical function in the hyporheic zone. <i>Science of the Total Environment</i> , 2018, 642, 742-753.	3.9	60
62	Dispersal limitation and thermodynamic constraints govern spatial structure of permafrost microbial communities. <i>FEMS Microbiology Ecology</i> , 2018, 94, .	1.3	62
63	Nearly a decade-long repeatable seasonal diversity patterns of bacterioplankton communities in the eutrophic Lake Donghu (Wuhan, China). <i>Molecular Ecology</i> , 2017, 26, 3839-3850.	2.0	76
64	Colonization Habitat Controls Biomass, Composition, and Metabolic Activity of Attached Microbial Communities in the Columbia River Hyporheic Corridor. <i>Applied and Environmental Microbiology</i> , 2017, 83, .	1.4	20
65	Autogenic succession and deterministic recovery following disturbance in soil bacterial communities. <i>Scientific Reports</i> , 2017, 7, 45691.	1.6	71
66	Deterministic influences exceed dispersal effects on hydrologically-connected microbiomes. <i>Environmental Microbiology</i> , 2017, 19, 1552-1567.	1.8	143
67	Geochemical and Microbial Community Attributes in Relation to Hyporheic Zone Geological Facies. <i>Scientific Reports</i> , 2017, 7, 12006.	1.6	40
68	Long-term nitrogen addition affects the phylogenetic turnover of soil microbial community responding to moisture pulse. <i>Scientific Reports</i> , 2017, 7, 17492.	1.6	79
69	Dispersal-Based Microbial Community Assembly Decreases Biogeochemical Function. <i>Processes</i> , 2017, 5, 65.	1.3	93
70	Regulation-Structured Dynamic Metabolic Model Provides a Potential Mechanism for Delayed Enzyme Response in Denitrification Process. <i>Frontiers in Microbiology</i> , 2017, 8, 1866.	1.5	40
71	Soil respiration across a permafrost transition zone: spatial structure and environmental correlates. <i>Biogeosciences</i> , 2017, 14, 4341-4354.	1.3	7
72	Biogeochemical cycling at the aquatic-terrestrial interface is linked to parafluvial hyporheic zone inundation history. <i>Biogeosciences</i> , 2017, 14, 4229-4241.	1.3	25

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73	Carbon Inputs From Riparian Vegetation Limit Oxidation of Physically Bound Organic Carbon Via Biochemical and Thermodynamic Processes. <i>Journal of Geophysical Research G: Biogeosciences</i> , 2017, 122, 3188-3205.	1.3	58
74	Aligning the Measurement of Microbial Diversity with Macroecological Theory. <i>Frontiers in Microbiology</i> , 2016, 7, 1487.	1.5	13
75	Coupling Spatiotemporal Community Assembly Processes to Changes in Microbial Metabolism. <i>Frontiers in Microbiology</i> , 2016, 7, 1949.	1.5	87
76	Spatial and successional dynamics of microbial biofilm communities in a grassland stream ecosystem. <i>Molecular Ecology</i> , 2016, 25, 4674-4688.	2.0	59
77	Seasonal hyporheic dynamics control coupled microbiology and geochemistry in Colorado River sediments. <i>Journal of Geophysical Research G: Biogeosciences</i> , 2016, 121, 2976-2987.	1.3	49
78	Coupling among Microbial Communities, Biogeochemistry and Mineralogy across Biogeochemical Facies. <i>Scientific Reports</i> , 2016, 6, 30553.	1.6	26
79	Groundwater-surface water mixing shifts ecological assembly processes and stimulates organic carbon turnover. <i>Nature Communications</i> , 2016, 7, 11237.	5.8	290
80	Estimating and mapping ecological processes influencing microbial community assembly. <i>Frontiers in Microbiology</i> , 2015, 6, 370.	1.5	578
81	The reduced genomes of Parcubacteria (OD1) contain signatures of a symbiotic lifestyle. <i>Frontiers in Microbiology</i> , 2015, 6, 713.	1.5	280
82	Relative Roles of Deterministic and Stochastic Processes in Driving the Vertical Distribution of Bacterial Communities in a Permafrost Core from the Qinghai-Tibet Plateau, China. <i>PLoS ONE</i> , 2015, 10, e0145747.	1.1	44
83	Disentangling mechanisms that mediate the balance between stochastic and deterministic processes in microbial succession. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, E1326-32.	3.3	972
84	The Gut Microbiota of Rural Papua New Guineans: Composition, Diversity Patterns, and Ecological Processes. <i>Cell Reports</i> , 2015, 11, 527-538.	2.9	475
85	On the processes generating latitudinal richness gradients: identifying diagnostic patterns and predictions. <i>Frontiers in Genetics</i> , 2014, 5, 420.	1.1	27
86	When should species richness be energy limited, and how would we know?. <i>Ecology Letters</i> , 2014, 17, 401-413.	3.0	107
87	Linking microbial community structure to β -glucosidic function in soil aggregates. <i>ISME Journal</i> , 2013, 7, 2044-2053.	4.4	110
88	Quantifying community assembly processes and identifying features that impose them. <i>ISME Journal</i> , 2013, 7, 2069-2079.	4.4	1,354
89	Stochastic and deterministic drivers of spatial and temporal turnover in breeding bird communities. <i>Global Ecology and Biogeography</i> , 2013, 22, 202-212.	2.7	121
90	Phylogenetic beta diversity in bacterial assemblages across ecosystems: deterministic versus stochastic processes. <i>ISME Journal</i> , 2013, 7, 1310-1321.	4.4	515

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91	An empirical assessment of tree branching networks and implications for plant allometric scaling models. <i>Ecology Letters</i> , 2013, 16, 1069-1078.	3.0	89
92	Correlations between physical and chemical defences in plants: tradeoffs, syndromes, or just many different ways to skin a herbivorous cat?. <i>New Phytologist</i> , 2013, 198, 252-263.	3.5	124
93	The epsomitic phototrophic microbial mat of Hot Lake, Washington: community structural responses to seasonal cycling. <i>Frontiers in Microbiology</i> , 2013, 4, 323.	1.5	75
94	Evolving ecological networks and the emergence of biodiversity patterns across temperature gradients. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2012, 279, 1051-1060.	1.2	40
95	Dispersal, environmental niches and oceanic-scale turnover in deep-sea bivalves. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2012, 279, 1993-2002.	1.2	54
96	Interannual variability of growth and reproduction in <i>Bursera simaruba</i> : the role of allometry and resource variability. <i>Ecology</i> , 2012, 93, 180-190.	1.5	19
97	Response to Comments on "Disentangling the Drivers of β^2 Diversity Along Latitudinal and Elevational Gradients". <i>Science</i> , 2012, 335, 1573-1573.	6.0	8
98	Eco-Evolutionary Community Dynamics: Covariation between Diversity and Invasibility across Temperature Gradients. <i>American Naturalist</i> , 2012, 180, E110-E126.	1.0	9
99	Testing the metabolic theory of ecology. <i>Ecology Letters</i> , 2012, 15, 1465-1474.	3.0	155
100	Stochastic and deterministic assembly processes in subsurface microbial communities. <i>ISME Journal</i> , 2012, 6, 1653-1664.	4.4	1,203
101	Temporal turnover in the composition of tropical tree communities: functional determinism and phylogenetic stochasticity. <i>Ecology</i> , 2012, 93, 490-499.	1.5	168
102	The biogeography and filtering of woody plant functional diversity in North and South America. <i>Global Ecology and Biogeography</i> , 2012, 21, 798-808.	2.7	235
103	Navigating the multiple meanings of β^2 diversity: a roadmap for the practicing ecologist. <i>Ecology Letters</i> , 2011, 14, 19-28.	3.0	1,899
104	Variation in above-ground forest biomass across broad climatic gradients. <i>Global Ecology and Biogeography</i> , 2011, 20, 744-754.	2.7	195
105	Putting plant resistance traits on the map: a test of the idea that plants are better defended at lower latitudes. <i>New Phytologist</i> , 2011, 191, 777-788.	3.5	155
106	Disentangling the Drivers of β^2 Diversity Along Latitudinal and Elevational Gradients. <i>Science</i> , 2011, 333, 1755-1758.	6.0	617
107	Trophic ecology of an aquatic mite (<i>Piona carnea</i>) preying on <i>Daphnia pulex</i> : effects of predator density, nutrient supply and a second predator (<i>Chaoborus americanus</i>). <i>Hydrobiologia</i> , 2011, 668, 171-182.	1.0	1
108	Inferring Ecological Processes from Taxonomic, Phylogenetic and Functional Trait β^2 -Diversity. <i>PLoS ONE</i> , 2011, 6, e20906.	1.1	69

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109	Integrating elements and energy through the metabolic dependencies of gross growth efficiency and the threshold elemental ratio. <i>Oikos</i> , 2010, 119, 752.	1.2	0
110	Integrating elements and energy through the metabolic dependencies of gross growth efficiency and the threshold elemental ratio. <i>Oikos</i> , 2010, 119, 752-765.	1.2	51
111	Functional trait assembly through ecological and evolutionary time. <i>Theoretical Ecology</i> , 2009, 2, 239-250.	0.4	19
112	Advancing the metabolic theory of biodiversity. <i>Ecology Letters</i> , 2009, 12, 1001-1015.	3.0	68
113	Above-ground forest biomass is not consistently related to wood density in tropical forests. <i>Global Ecology and Biogeography</i> , 2009, 18, 617-625.	2.7	46
114	On the relationship between mass and diameter distributions in tree communities. <i>Ecology Letters</i> , 2008, 11, 1287-1293.	3.0	13
115	The control of color change in the Pacific tree frog, <i>Hyla regilla</i> . <i>Canadian Journal of Zoology</i> , 2004, 82, 889-896.	0.4	39