

# John M Eppley

## List of Publications by Year in descending order

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Version: 2024-02-01

27  
papers

2,599  
citations

394421

19  
h-index

552781

26  
g-index

33  
all docs

33  
docs citations

33  
times ranked

2959  
citing authors

#	ARTICLE	IF	CITATIONS
1	Community-led, integrated, reproducible multi-omics with anvii™. <i>Nature Microbiology</i> , 2021, 6, 3-6.	13.3	370
2	Multispecies diel transcriptional oscillations in open ocean heterotrophic bacterial assemblages. <i>Science</i> , 2014, 345, 207-212.	12.6	245
3	Microbial community transcriptional networks are conserved in three domains at ocean basin scales. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 5443-5448.	7.1	225
4	Environmental drivers of a microbial genomic transition zone in the ocean's interior. <i>Nature Microbiology</i> , 2017, 2, 1367-1373.	13.3	177
5	Pattern and synchrony of gene expression among sympatric marine microbial populations. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, E488-97.	7.1	164
6	Biological composition and microbial dynamics of sinking particulate organic matter at abyssal depths in the oligotrophic open ocean. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 11824-11832.	7.1	150
7	Microbial community structure and function on sinking particles in the North Pacific Subtropical Gyre. <i>Frontiers in Microbiology</i> , 2015, 6, 469.	3.5	148
8	Planktonic Euryarchaeota are a significant source of archaeal tetraether lipids in the ocean. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 9858-9863.	7.1	134
9	Microbial community phylogenetic and trait diversity declines with depth in a marine oxygen minimum zone. <i>Ecology</i> , 2012, 93, 1659-1673.	3.2	129
10	Coordinated regulation of growth, activity and transcription in natural populations of the unicellular nitrogen-fixing cyanobacterium <i>Crocospaera</i> . <i>Nature Microbiology</i> , 2017, 2, 17118.	13.3	122
11	Diel cycling and long-term persistence of viruses in the ocean's euphotic zone. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 11446-11451.	7.1	116
12	Assembly-free single-molecule sequencing recovers complete virus genomes from natural microbial communities. <i>Genome Research</i> , 2020, 30, 437-446.	5.5	80
13	Wind and sunlight shape microbial diversity in surface waters of the North Pacific Subtropical Gyre. <i>ISME Journal</i> , 2016, 10, 1308-1322.	9.8	73
14	Genetic Exchange Across a Species Boundary in the Archaeal Genus <i>Ferroplasma</i> . <i>Genetics</i> , 2007, 177, 407-416.	2.9	67
15	Double-stranded DNA viroplankton dynamics and reproductive strategies in the oligotrophic open ocean water column. <i>ISME Journal</i> , 2020, 14, 1304-1315.	9.8	65
16	Microbial dynamics of elevated carbon flux in the open ocean's abyss. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	65
17	Draft genome sequence of marine alphaproteobacterial strain HIMB11, the first cultivated representative of a unique lineage within the Roseobacter clade possessing an unusually small genome. <i>Standards in Genomic Sciences</i> , 2014, 9, 632-645.	1.5	40
18	Time-series analyses of Monterey Bay coastal microbial picoplankton using a genome proxy™ microarray. <i>Environmental Microbiology</i> , 2011, 13, 116-134.	3.8	31

#	ARTICLE	IF	CITATIONS
19	Metapangenomics reveals depth-dependent shifts in metabolic potential for the ubiquitous marine bacterial SAR324 lineage. <i>Microbiome</i> , 2021, 9, 172.	11.1	25
20	Improved Environmental Genomes via Integration of Metagenomic and Single-Cell Assemblies. <i>Frontiers in Microbiology</i> , 2016, 7, 143.	3.5	24
21	Complex marine microbial communities partition metabolism of scarce resources over the diel cycle. <i>Nature Ecology and Evolution</i> , 2022, 6, 218-229.	7.8	21
22	Diverse Genomic Traits Differentiate Sinking-Particle-Associated versus Free-Living Microbes throughout the Oligotrophic Open Ocean Water Column. <i>MBio</i> , 2022, 13, .	4.1	21
23	Reply to Schouten et al.: Marine Group II planktonic Euryarchaeota are significant contributors to tetraether lipids in the ocean. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, E4286.	7.1	20
24	Diel Oscillation of Microbial Gene Transcripts Declines With Depth in Oligotrophic Ocean Waters. <i>Frontiers in Microbiology</i> , 2019, 10, 2191.	3.5	19
25	Diversity and origins of bacterial and archaeal viruses on sinking particles reaching the abyssal ocean. <i>ISME Journal</i> , 2022, 16, 1627-1635.	9.8	18
26	Microbial Sources of Exocellular DNA in the Ocean. <i>Applied and Environmental Microbiology</i> , 2022, 88, e0209321.	3.1	6
27	Novel Integrative Elements and Genomic Plasticity in Ocean Ecosystems. <i>SSRN Electronic Journal</i> , 0, , .	0.4	2