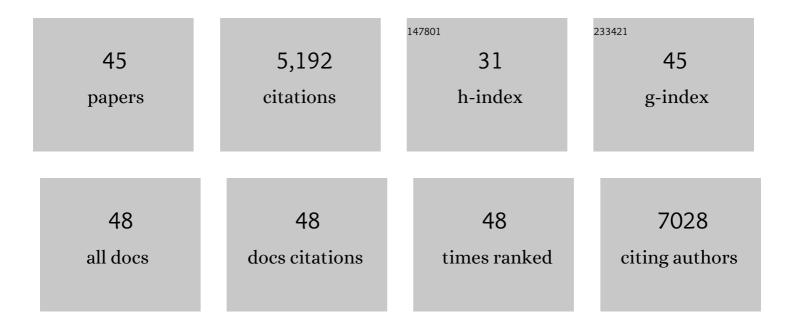
## Nicole J Poulton

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8626932/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Marine DNA Viral Macro- and Microdiversity from Pole to Pole. Cell, 2019, 177, 1109-1123.e14.	28.9	541
2	Potential for Chemolithoautotrophy Among Ubiquitous Bacteria Lineages in the Dark Ocean. Science, 2011, 333, 1296-1300.	12.6	510
3	Prevalent genome streamlining and latitudinal divergence of planktonic bacteria in the surface ocean. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 11463-11468.	7.1	328
4	Global Trends in Marine Plankton Diversity across Kingdoms of Life. Cell, 2019, 179, 1084-1097.e21.	28.9	271
5	Gene Expression Changes and Community Turnover Differentially Shape the Global Ocean Metatranscriptome. Cell, 2019, 179, 1068-1083.e21.	28.9	268
6	Obtaining genomes from uncultivated environmental microorganisms using FACS–based single-cell genomics. Nature Protocols, 2014, 9, 1038-1048.	12.0	243
7	Capturing Single Cell Genomes of Active Polysaccharide Degraders: An Unexpected Contribution of Verrucomicrobia. PLoS ONE, 2012, 7, e35314.	2.5	236
8	Phylogenetic Diversity and Specificity of Bacteria Closely Associated with Alexandrium spp. and Other Phytoplankton. Applied and Environmental Microbiology, 2005, 71, 3483-3494.	3.1	198
9	The ocean sampling day consortium. GigaScience, 2015, 4, 27.	6.4	185
10	Accumulation and enhanced cycling of polyphosphate by Sargasso Sea plankton in response to low phosphorus. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 8089-8094.	7.1	172
11	Improved genome recovery and integrated cell-size analyses of individual uncultured microbial cells and viral particles. Nature Communications, 2017, 8, 84.	12.8	169
12	High-throughput single-cell sequencing identifies photoheterotrophs and chemoautotrophs in freshwater bacterioplankton. ISME Journal, 2012, 6, 113-123.	9.8	168
13	Charting the Complexity of the Marine Microbiome through Single-Cell Genomics. Cell, 2019, 179, 1623-1635.e11.	28.9	158
14	Viral to metazoan marine plankton nucleotide sequences from the Tara Oceans expedition. Scientific Data, 2017, 4, 170093.	5.3	147
15	Rokubacteria: Genomic Giants among the Uncultured Bacterial Phyla. Frontiers in Microbiology, 2017, 8, 2264.	3.5	142
16	Unveiling <i>in situ</i> interactions between marine protists and bacteria through single cell sequencing. ISME Journal, 2012, 6, 703-707.	9.8	124
17	Cryptic and abundant marine viruses at the evolutionary origins of Earth's RNA virome. Science, 2022, 376, 156-162.	12.6	124
18	Single-cell genomics of multiple uncultured stramenopiles reveals underestimated functional diversity across oceans. Nature Communications, 2018, 9, 310.	12.8	101

NICOLE J POULTON

#	Article	IF	CITATIONS
19	Genomic and Metabolic Diversity of Marine Group I Thaumarchaeota in the Mesopelagic of Two Subtropical Gyres. PLoS ONE, 2014, 9, e95380.	2.5	95
20	Distribution of planktonic aerobic anoxygenic photoheterotrophic bacteria in the northwest Atlantic. Limnology and Oceanography, 2006, 51, 38-46.	3.1	93
21	Capturing diversity of marine heterotrophic protists: one cell at a time. ISME Journal, 2011, 5, 674-684.	9.8	86
22	What's New Is Old: Resolving the Identity of Leptothrix ochracea Using Single Cell Genomics, Pyrosequencing and FISH. PLoS ONE, 2011, 6, e17769.	2.5	85
23	Counting heterotrophic nanoplanktonic protists in cultures and aquatic communities by flow cytometry. Aquatic Microbial Ecology, 2004, 34, 263-277.	1.8	84
24	A simple optical index shows spatial and temporal heterogeneity in phytoplankton community composition during the 2008 North Atlantic Bloom Experiment. Biogeosciences, 2015, 12, 2179-2194.	3.3	78
25	Imaging flow cytometry for phytoplankton analysis. Methods, 2017, 112, 188-200.	3.8	76
26	Single cell genome analysis supports a link between phagotrophy and primary plastid endosymbiosis. Scientific Reports, 2012, 2, 356.	3.3	62
27	Ancestral Absence of Electron Transport Chains in Patescibacteria and DPANN. Frontiers in Microbiology, 2020, 11, 1848.	3.5	62
28	Compendium of 530 metagenome-assembled bacterial and archaeal genomes from the polar Arctic Ocean. Nature Microbiology, 2021, 6, 1561-1574.	13.3	57
29	Toxin variability in natural populations of Alexandrium fundyense in Casco Bay, Maine—evidence of nitrogen limitation. Deep-Sea Research Part II: Topical Studies in Oceanography, 2005, 52, 2501-2521.	1.4	40
30	Genomic exploration of individual giant ocean viruses. ISME Journal, 2017, 11, 1736-1745.	9.8	40
31	Anatomy of a red tide bloom off the southwest coast of Florida. Harmful Algae, 2008, 7, 817-826.	4.8	36
32	FlowCam: Quantification and Classification of Phytoplankton by Imaging Flow Cytometry. Methods in Molecular Biology, 2016, 1389, 237-247.	0.9	32
33	Single Cell Genomics Reveals Viruses Consumed by Marine Protists. Frontiers in Microbiology, 2020, 11, 524828.	3.5	26
34	Automated Isolation Techniques for Microalgae. , 2005, , 101-116.		26
35	Targeted Sorting of Single Virus-Infected Cells of the Coccolithophore Emiliania huxleyi. PLoS ONE, 2011, 6, e22520.	2.5	23
36	Temporal dynamics of eukaryotic microbial diversity at a coastal Pacific site. ISME Journal, 2018, 12, 2278-2291.	9.8	19

NICOLE J POULTON

#	Article	IF	CITATIONS
37	Characterizing the phytoplankton soup: pump and plumbing effects on the particle assemblage in underway optical seawater systems. Optics Express, 2016, 24, 20703.	3.4	18
38	Best practices in plant cytometry. Cytometry Part A: the Journal of the International Society for Analytical Cytology, 2021, 99, 311-317.	1.5	16
39	Radiometric approach for the detection of picophytoplankton assemblages across oceanic fronts. Optics Express, 2020, 28, 25682.	3.4	12
40	Single cell ecogenomics reveals mating types of individual cells and ssDNA viral infections in the smallest photosynthetic eukaryotes. Philosophical Transactions of the Royal Society B: Biological Sciences, 2019, 374, 20190089.	4.0	11
41	Zooplankton grazing and egestion shifts particle size distribution in natural communities. Marine Ecology - Progress Series, 2017, 575, 43-56.	1.9	8
42	Lighting up phytoplankton cells with quantum dots. Limnology and Oceanography: Methods, 2008, 6, 653-658.	2.0	4
43	Inter-laboratory validation of the serial dilution culture—most probable number method for enumerating viable phytoplankton. Journal of Applied Phycology, 2019, 31, 491-503.	2.8	3
44	Genome to phenome tools: In vivo and in vitro transfection of Crassostrea virginica hemocytes. Fish and Shellfish Immunology, 2020, 103, 438-441.	3.6	3
45	Oceanic Crustal Fluid Single Cell Genomics Complements Metagenomic and Metatranscriptomic Surveys With Orders of Magnitude Less Sample Volume. Frontiers in Microbiology, 2021, 12, 738231.	3.5	1