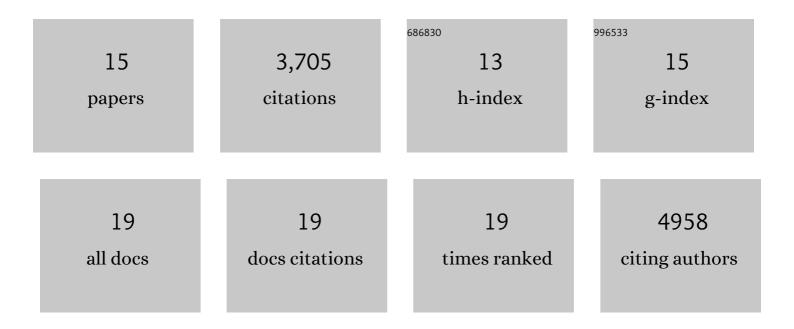
Ann C Gregory

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/10639008/publications.pdf Version: 2024-02-01



ANN C CRECORY

#	Article	IF	CITATIONS
1	Microbiota Transfer Therapy alters gut ecosystem and improves gastrointestinal and autism symptoms: an open-label study. Microbiome, 2017, 5, 10.	4.9	901
2	Patterns and ecological drivers of ocean viral communities. Science, 2015, 348, 1261498.	6.0	617
3	Marine DNA Viral Macro- and Microdiversity from Pole to Pole. Cell, 2019, 177, 1109-1123.e14.	13.5	541
4	The Gut Virome Database Reveals Age-Dependent Patterns of Virome Diversity in the Human Gut. Cell Host and Microbe, 2020, 28, 724-740.e8.	5.1	352
5	Global Trends in Marine Plankton Diversity across Kingdoms of Life. Cell, 2019, 179, 1084-1097.e21.	13.5	271
6	Gene Expression Changes and Community Turnover Differentially Shape the Global Ocean Metatranscriptome. Cell, 2019, 179, 1068-1083.e21.	13.5	268
7	Viral tagging reveals discrete populations in Synechococcus viral genome sequence space. Nature, 2014, 513, 242-245.	13.7	183
8	Synthetic ecology of the human gut microbiota. Nature Reviews Microbiology, 2019, 17, 754-763.	13.6	117
9	Long-read viral metagenomics captures abundant and microdiverse viral populations and their niche-defining genomic islands. PeerJ, 2019, 7, e6800.	0.9	109
10	Genomic differentiation among wild cyanophages despite widespread horizontal gene transfer. BMC Genomics, 2016, 17, 930.	1.2	96
11	Potential virus-mediated nitrogen cycling in oxygen-depleted oceanic waters. ISME Journal, 2021, 15, 981-998.	4.4	71
12	Clinical practices underlie COVID-19 patient respiratory microbiome composition and its interactions with the host. Nature Communications, 2021, 12, 6243.	5.8	42
13	Smoking is associated with quantifiable differences in the human lung DNA virome and metabolome. Respiratory Research, 2018, 19, 174.	1.4	28
14	MetaPop: a pipeline for macro- and microdiversity analyses and visualization of microbial and viral metagenome-derived populations. Microbiome, 2022, 10, 49.	4.9	24
15	efam: an <i>e</i> xpanded, metaproteome-supported HMM profile database of viral protein <i>fam</i> ilies. Bioinformatics, 2021, 37, 4202-4208.	1.8	15