Ann C Gregory

List of Publications by Year in descending order

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Version: 2024-02-01

686830 996533 3,705 15 13 15 citations h-index g-index papers 19 19 19 4958 docs citations times ranked citing authors all docs

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