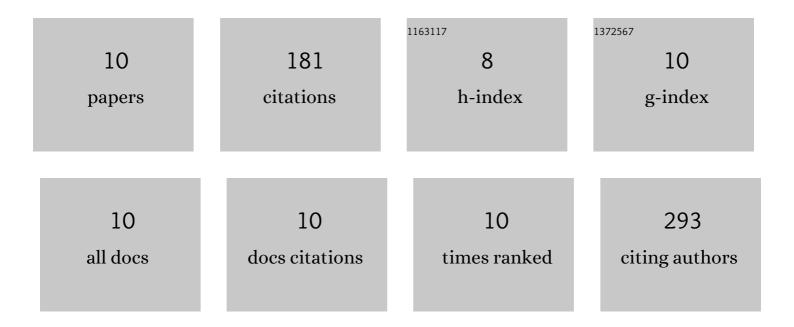
Katterinne N Mendez

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1983076/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	In-Depth Genomic and Phenotypic Characterization of the Antarctic Psychrotolerant Strain Pseudomonas sp. MPC6 Reveals Unique Metabolic Features, Plasticity, and Biotechnological Potential. Frontiers in Microbiology, 2019, 10, 1154.	3.5	36
2	Comparative genomic analysis of a new tellurite-resistant <i>Psychrobacter</i> strain isolated from the Antarctic Peninsula. PeerJ, 2018, 6, e4402.	2.0	30
3	Genomic Analysis of 48 Paenibacillus larvae Bacteriophages. Viruses, 2018, 10, 377.	3.3	26
4	Variability in Genomic and Virulent Properties of Porphyromonas gingivalis Strains Isolated From Healthy and Severe Chronic Periodontitis Individuals. Frontiers in Cellular and Infection Microbiology, 2019, 9, 246.	3.9	19
5	Copper-induced increased expression of genes involved in photosynthesis, carotenoid synthesis and C assimilation in the marine alga Ulva compressa. BMC Genomics, 2018, 19, 829.	2.8	18
6	Living to the High Extreme: Unraveling the Composition, Structure, and Functional Insights of Bacterial Communities Thriving in the Arsenic-Rich Salar de Huasco Altiplanic Ecosystem. Microbiology Spectrum, 2021, 9, e0044421.	3.0	16
7	Genomic Variation and Arsenic Tolerance Emerged as Niche Specific Adaptations by Different Exiguobacterium Strains Isolated From the Extreme Salar de Huasco Environment in Chilean – Altiplano. Frontiers in Microbiology, 2020, 11, 1632.	3.5	14
8	Complete genome sequence of the marine Rhodococcus sp. H-CA8f isolated from Comau fjord in Northern Patagonia, Chile. Marine Genomics, 2018, 40, 13-17.	1.1	9
9	RNA-seq analysis of compensatory growth in the skeletal muscle of fine flounder (Paralichthys) Tj ETQq1 1 0.784	314 rgBT , 3.5	/Oyerlock 10 _
10	Computational Methods for Human Microbiome Analysis. Current Protocols in Microbiology, 2017, 47, 1E.14.1-1E.14.17.	6.5	4