

Katterinne N Mendez

List of Publications by Year in descending order

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Version: 2024-02-01

10
papers

181
citations

1163117

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h-index

1372567

10
g-index

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all docs

10
docs citations

10
times ranked

293
citing authors

#	ARTICLE	IF	CITATIONS
1	In-Depth Genomic and Phenotypic Characterization of the Antarctic Psychrotolerant Strain <i>Pseudomonas</i> sp. MPC6 Reveals Unique Metabolic Features, Plasticity, and Biotechnological Potential. <i>Frontiers in Microbiology</i> , 2019, 10, 1154.	3.5	36
2	Comparative genomic analysis of a new tellurite-resistant <i>Psychrobacter</i> strain isolated from the Antarctic Peninsula. <i>PeerJ</i> , 2018, 6, e4402.	2.0	30
3	Genomic Analysis of 48 <i>Paenibacillus</i> larvae Bacteriophages. <i>Viruses</i> , 2018, 10, 377.	3.3	26
4	Variability in Genomic and Virulent Properties of <i>Porphyromonas gingivalis</i> Strains Isolated From Healthy and Severe Chronic Periodontitis Individuals. <i>Frontiers in Cellular and Infection Microbiology</i> , 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 <i>Ulva compressa</i> . <i>BMC Genomics</i> , 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. <i>Microbiology Spectrum</i> , 2021, 9, e0044421.	3.0	16
7	Genomic Variation and Arsenic Tolerance Emerged as Niche Specific Adaptations by Different <i>Exiguobacterium</i> Strains Isolated From the Extreme Salar de Huasco Environment in Chilean “Altiplano”. <i>Frontiers in Microbiology</i> , 2020, 11, 1632.	3.5	14
8	Complete genome sequence of the marine <i>Rhodococcus</i> sp. H-CA8f isolated from Comau fjord in Northern Patagonia, Chile. <i>Marine Genomics</i> , 2018, 40, 13-17.	1.1	9
9	RNA-seq analysis of compensatory growth in the skeletal muscle of fine flounder (<i>Paralichthys</i>) Tj ETQq1 1 0.784314 rgBT /Overlock 1	3.5	9
10	Computational Methods for Human Microbiome Analysis. <i>Current Protocols in Microbiology</i> , 2017, 47, 1E.14.1-1E.14.17.	6.5	4