

# Katterinne N Mendez

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1983076/publications.pdf>

Version: 2024-02-01

10  
papers

181  
citations

1163117

8  
h-index

1372567

10  
g-index

10  
all docs

10  
docs citations

10  
times ranked

293  
citing authors

#	ARTICLE	IF	CITATIONS
1	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
2	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". <i>Frontiers in Microbiology</i> , 2020, 11, 1632.	3.5	14
3	Variability in Genomic and Virulent Properties of Porphyromonas gingivalis Strains Isolated From Healthy and Severe Chronic Periodontitis Individuals. <i>Frontiers in Cellular and Infection Microbiology</i> , 2019, 9, 246.	3.9	19
4	In-Depth Genomic and Phenotypic Characterization of the Antarctic Psychrotolerant Strain Pseudomonas sp. MPC6 Reveals Unique Metabolic Features, Plasticity, and Biotechnological Potential. <i>Frontiers in Microbiology</i> , 2019, 10, 1154.	3.5	36
5	Complete genome sequence of the marine Rhodococcus sp. H-CA8f isolated from Comau fjord in Northern Patagonia, Chile. <i>Marine Genomics</i> , 2018, 40, 13-17.	1.1	9
6	RNA-seq analysis of compensatory growth in the skeletal muscle of fine flounder (Paralichthys Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 54	3.5	9
7	Copper-induced increased expression of genes involved in photosynthesis, carotenoid synthesis and C assimilation in the marine alga Ulva compressa. <i>BMC Genomics</i> , 2018, 19, 829.	2.8	18
8	Genomic Analysis of 48 Paenibacillus larvae Bacteriophages. <i>Viruses</i> , 2018, 10, 377.	3.3	26
9	Comparative genomic analysis of a new tellurite-resistant Psychrobacter strain isolated from the Antarctic Peninsula. <i>PeerJ</i> , 2018, 6, e4402.	2.0	30
10	Computational Methods for Human Microbiome Analysis. <i>Current Protocols in Microbiology</i> , 2017, 47, 1E.14.1-1E.14.17.	6.5	4