

# Paola Bertolino

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2831125/publications.pdf>

Version: 2024-02-01

11  
papers

513  
citations

1040056

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1281871

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docs citations

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times ranked

695  
citing authors

#	ARTICLE	IF	CITATIONS
1	Environmental drivers of plankton protist communities along latitudinal and vertical gradients in the oldest and deepest freshwater lake. <i>Environmental Microbiology</i> , 2021, 23, 1436-1451.	3.8	22
2	Small freshwater ecosystems with dissimilar microbial communities exhibit similar temporal patterns. <i>Molecular Ecology</i> , 2021, 30, 2162-2177.	3.9	15
3	Rapid formation of mature microbialites in Lake Alchichica, Mexico. <i>Environmental Microbiology Reports</i> , 2021, 13, 600-605.	2.4	2
4	Core microbial communities of lacustrine microbialites sampled along an alkalinity gradient. <i>Environmental Microbiology</i> , 2021, 23, 51-68.	3.8	26
5	Hyperdiverse archaea near life limits at the polyextreme geothermal Dallol area. <i>Nature Ecology and Evolution</i> , 2019, 3, 1552-1561.	7.8	62
6	Unveiling microbial interactions in stratified mat communities from a warm saline shallow pond. <i>Environmental Microbiology</i> , 2017, 19, 2405-2421.	3.8	35
7	Resilience of Freshwater Communities of Small Microbial Eukaryotes Undergoing Severe Drought Events. <i>Frontiers in Microbiology</i> , 2016, 7, 812.	3.5	26
8	Comparative metagenomics unveils functions and genome features of microbialite-associated communities along a depth gradient. <i>Environmental Microbiology</i> , 2016, 18, 4990-5004.	3.8	30
9	Metagenome-based diversity analyses suggest a significant contribution of non-cyanobacterial lineages to carbonate precipitation in modern microbialites. <i>Frontiers in Microbiology</i> , 2015, 6, 797.	3.5	50
10	Marked seasonality and high spatial variability of protist communities in shallow freshwater systems. <i>ISME Journal</i> , 2015, 9, 1941-1953.	9.8	165
11	Complex communities of small protists and unexpected occurrence of typical marine lineages in shallow freshwater systems. <i>Environmental Microbiology</i> , 2015, 17, 3610-3627.	3.8	80