List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Microbiology of the Red Sea (and other) deepâ€sea anoxic brine lakes. Environmental Microbiology Reports, 2011, 3, 416-433.	1.0	158
2	Halorhabdus tiamatea sp. nov., a non-pigmented, extremely halophilic archaeon from a deep-sea, hypersaline anoxic basin of the Red Sea, and emended description of the genus Halorhabdus. International Journal of Systematic and Evolutionary Microbiology, 2008, 58, 215-220.	0.8	124
3	Rhizosphere microbiome metagenomics of gray mangroves (Avicennia marina) in the Red Sea. Gene, 2016, 576, 626-636.	1.0	116
4	Biogeography of pelagic bacterioplankton across an antagonistic temperature–salinity gradient in the Red Sea. Molecular Ecology, 2012, 21, 388-405.	2.0	98
5	A New Lineage of Halophilic, Wall-Less, Contractile Bacteria from a Brine-Filled Deep of the Red Sea. Journal of Bacteriology, 2008, 190, 3580-3587.	1.0	84
6	INDIGO – INtegrated Data Warehouse of Mlcrobial GenOmes with Examples from the Red Sea Extremophiles. PLoS ONE, 2013, 8, e82210.	1.1	83
7	Description of Idiomarina insulisalsae sp. nov., isolated from the soil of a sea salt evaporation pond, proposal to transfer the species of the genus Pseudidiomarina to the genus Idiomarina and emended description of the genus Idiomarina. Systematic and Applied Microbiology, 2009, 32, 371-378.	1.2	77
8	Salinisphaera shabanensis gen. nov., sp. nov., a novel, moderately halophilic bacterium from the brine–seawater interface of the Shaban Deep, Red Sea. Extremophiles, 2003, 7, 29-34.	0.9	72
9	Soil and Rhizosphere Associated Fungi in Gray Mangroves (Avicennia marina) from the Red Sea — A Metagenomic Approach. Genomics, Proteomics and Bioinformatics, 2015, 13, 310-320.	3.0	67
10	Marinobacter salsuginis sp. nov., isolated from the brine–seawater interface of the Shaban Deep, Red Sea. International Journal of Systematic and Evolutionary Microbiology, 2007, 57, 1035-1040.	0.8	61
11	Comparative genomics reveals adaptations of a halotolerant thaumarchaeon in the interfaces of brine pools in the Red Sea. ISME Journal, 2015, 9, 396-411.	4.4	60
12	Leuconostoc ficulneum sp. nov., a novel lactic acid bacterium isolated from a ripe fig, and reclassification of Lactobacillus fructosus as Leuconostoc fructosum comb. nov International Journal of Systematic and Evolutionary Microbiology, 2002, 52, 647-655.	0.8	57
13	Unique Prokaryotic Consortia in Geochemically Distinct Sediments from Red Sea Atlantis II and Discovery Deep Brine Pools. PLoS ONE, 2012, 7, e42872.	1.1	45
14	Diversity of methanogens and sulfate-reducing bacteria in the interfaces of five deep-sea anoxic brines of the Red Sea. Research in Microbiology, 2015, 166, 688-699.	1.0	43
15	<scp><i>H</i></scp> <i>alorhabdus tiamatea:</i> proteogenomics and glycosidase activity measurements identify the first cultivated euryarchaeon from a deepâ€sea anoxic brine lake as potential polysaccharide degrader. Environmental Microbiology, 2014, 16, 2525-2537.	1.8	41
16	First Insights into the Viral Communities of the Deep-sea Anoxic Brines of the Red Sea. Genomics, Proteomics and Bioinformatics, 2015, 13, 304-309.	3.0	33
17	Cytotoxic and apoptotic evaluations of marine bacteria isolated from brine-seawater interface of the Red Sea. BMC Complementary and Alternative Medicine, 2013, 13, 29.	3.7	30
18	Earth's Stratosphere and Microbial Life. Current Issues in Molecular Biology, 2020, 38, 197-244.	1.0	27

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19	Genome Sequence of Haloplasma contractile, an Unusual Contractile Bacterium from a Deep-Sea Anoxic Brine Lake. Journal of Bacteriology, 2011, 193, 4551-4552.	1.0	26
20	Visualizing the invisible: class excursions to ignite children's enthusiasm for microbes. Microbial Biotechnology, 2020, 13, 844-887.	2.0	26
21	Experimental and Simulation Efforts in the Astrobiological Exploration of Exooceans. Space Science Reviews, 2020, 216, 9.	3.7	25
22	Genome Sequence of Halorhabdus tiamatea, the First Archaeon Isolated from a Deep-Sea Anoxic Brine Lake. Journal of Bacteriology, 2011, 193, 4553-4554.	1.0	24
23	Biological Contamination Prevention for Outer Solar System Moons of Astrobiological Interest: What Do We Need to Know?. Astrobiology, 2019, 19, 951-974.	1.5	24
24	Genome Sequence of Salinisphaera shabanensis, a Gammaproteobacterium from the Harsh, Variable Environment of the Brine-Seawater Interface of the Shaban Deep in the Red Sea. Journal of Bacteriology, 2011, 193, 4555-4556.	1.0	21
25	Core Microbial Functional Activities in Ocean Environments Revealed by Global Metagenomic Profiling Analyses. PLoS ONE, 2014, 9, e97338.	1.1	20
26	Mars: new insights and unresolved questions. International Journal of Astrobiology, 2021, 20, 394-426.	0.9	19
27	Mycogenic Metal Nanoparticles for the Treatment of Mycobacterioses. Antibiotics, 2020, 9, 569.	1.5	18
28	Genome Sequence of <i>Pseudomonas</i> sp. Strain Chol1, a Model Organism for the Degradation of Bile Salts and Other Steroid Compounds. Genome Announcements, 2013, 1, .	0.8	17
29	In silico exploration of Red Sea Bacillus genomes for natural product biosynthetic gene clusters. BMC Genomics, 2018, 19, 382.	1.2	17
30	Biogenic Metal Nanoparticles: A New Approach to Detect Life on Mars?. Life, 2020, 10, 28.	1.1	17
31	Exploring Deep-Sea Brines as Potential Terrestrial Analogues of Oceans in the Icy Moons of the Outer Solar System. Current Issues in Molecular Biology, 2020, 38, 123-162.	1.0	16
32	Microbial Diversity and Biosignatures: An Icy Moons Perspective. Space Science Reviews, 2020, 216, 1.	3.7	14
33	Habitability Models for Astrobiology. Astrobiology, 2021, 21, 1017-1027.	1.5	13
34	DESM: portal for microbial knowledge exploration systems. Nucleic Acids Research, 2016, 44, D624-D633.	6.5	12
35	Microbial Pathogenicity in Space. Pathogens, 2021, 10, 450.	1.2	11
36	Bioprospecting Archaea: Focus on Extreme Halophiles. Topics in Biodiversity and Conservation, 2017, , 81-112.	0.3	10

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37	Progression in the bacterial load during the breeding season in nest boxes occupied by the Blue Tit and its potential impact on hatching or fledging success. Journal of Ornithology, 2018, 159, 1009-1017.	0.5	10
38	Systematics, functional morphology and distribution of a bivalve (<i>Apachecorbula muriatica</i>) Tj ETQq0 Biological Association of the United Kingdom, 2015, 95, 523-535.	0 0 rgBT /Ove 0.4	erlock 10 Tf 50 9
39	Zooplankton at deep Red Sea brine pools. Journal of Plankton Research, 2016, 38, 679-684.	0.8	8
40	Fueling the Bio-economy: European Culture Collections and Microbiology Education and Training. Trends in Microbiology, 2016, 24, 77-79.	3.5	8
41	Bacterial species richness at three stages of the breeding season in Cyanistes caeruleus (blue tit). Acta Oecologica, 2018, 92, 123-130.	0.5	6
42	Motilimonas cestriensis sp. nov., isolated from an inland brine spring in Northern England. International Journal of Systematic and Evolutionary Microbiology, 2019, 71, .	0.8	6
43	Surviving Mars: new insights into the persistence of facultative anaerobic microbes from analogue sites. International Journal of Astrobiology, 2022, 21, 110-127.	0.9	6
44	Habitability Models for Planetary Sciences. , 2021, 53, .		3
45	Earth's Stratosphere and Microbial Life. , 2020, , .		3
46	Sulfur Cycling as a Viable Metabolism under Simulated Noachian/Hesperian Chemistries. Life, 2022, 12, 523.	1.1	3
47	The Family Salinisphaeraceae. , 2014, , 591-596.		2
48	The Family Haloplasmataceae. , 2014, , 179-184.		2
49	Haloferax profundi sp. nov. and Haloferax marisrubri sp. nov., Isolated from the Discovery Deep Brine-Seawater Interface in the Red Sea. Microorganisms, 2020, 8, 1475.	1.6	1
50	Geochemistry and Life at the Interfaces of Brine-Filled Deeps in the Red Sea. Springer Oceanography, 2019, , 185-194.	0.2	1
51	Exploring Deep-Sea Brines as Potential Terrestrial Analogues of Oceans in the Icy Moons of the Outer Solar System. , 2020, , .		1
52	Microbes from Brine Systems with Fluctuating Salinity Can Thrive under Simulated Martian Chemical Conditions. Life, 2022, 12, 12.	1.1	1
53	Offer, demand, and needs in training and education: a study focusing on microbial culture collections within the MIRRI Consortium. New Biotechnology, 2014, 31, S155.	2.4	0
54	Isolation of extremophilic bacteria from microbial mats and the applicability of their amylases for bioethanol production from food waste. Energy Sources, Part A: Recovery, Utilization and Environmental Effects, 0, , 1-12.	1.2	0

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55	Halophiles: the salt of the Earth. Biochemist, 2017, 39, 18-21.	0.2	ο
56	Genomic trends on the biogenic CaCO3 production in the genus Bacillus. Access Microbiology, 2019, 1,	0.2	0
57	Biomineralization of microbes in natural and restored saltmarshes: a missing link in restoration efforts?. Access Microbiology, 2019, 1, .	0.2	0
58	Metabolic profiling and environmental characterisation of salterns in the islands of Cabo Verde. Access Microbiology, 2019, 1, .	0.2	0
59	Motilimonas cestriensis sp. nov., isolated from a Cheshire brine spring. Access Microbiology, 2019, 1, .	0.2	0
60	Out of This World: From the Bottom of the Red Sea to the Red Planet. Frontiers for Young Minds, 0, 8,	0.8	0