

Ramn Alberto Batista-Garca

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

27
papers

297
citations

10
h-index

16
g-index

28
ext. papers

438
ext. citations

5.2
avg, IF

3.2
L-index

#	Paper	IF	Citations
27	The Atacama Desert: A Biodiversity Hotspot and Not Just a Mineral-Rich Region.. <i>Frontiers in Microbiology</i> , 2022 , 13, 812842	5.7	1
26	Surviving in the Brine: A Multi-Omics Approach for Understanding the Physiology of the Halophile Fungus at Saturated NaCl Concentration.. <i>Frontiers in Microbiology</i> , 2022 , 13, 840408	5.7	0
25	Transcriptomic analysis of polyaromatic hydrocarbon degradation by the halophilic fungus <i>Aspergillus sydowii</i> at hypersaline conditions. <i>Environmental Microbiology</i> , 2021 , 23, 3435-3459	5.2	13
24	Osmolyte Signatures for the Protection of Cells under Halophilic Conditions and Osmotic Shock. <i>Journal of Fungi (Basel, Switzerland)</i> , 2021 , 7,	5.6	1
23	ROS-Scavenging Enzymes as an Antioxidant Response to High Concentration of Anthracene in the Liverwort <i>L. Plants</i> , 2021 , 10,	4.5	2
22	Tracking gene expression, metabolic profiles, and biochemical analysis in the halotolerant basidiomycetous yeast <i>Rhodotorula mucilaginosa</i> EXF-1630 during benzo[a]pyrene and phenanthrene biodegradation under hypersaline conditions. <i>Environmental Pollution</i> , 2021 , 271, 116358	9.3	6
21	Transcriptional profiling reveals conserved and species-specific plant defense responses during the interaction of <i>Physcomitrium patens</i> with <i>Botrytis cinerea</i> . <i>Plant Molecular Biology</i> , 2021 , 107, 365-385	4.6	3
20	Infection by : Understanding the Fungal-Bryophyte Interaction by Microscopy, Phenomics and RNA Sequencing. <i>Journal of Fungi (Basel, Switzerland)</i> , 2021 , 7,	5.6	1
19	Effects on Plants Colonized with <i>P. Karst</i> Strains Genetically Modified in , a Gene Coding for a Protein with Expansin-like Activity. <i>Plants</i> , 2021 , 10,	4.5	1
18	The Microbial Composition in Circumneutral Thermal Springs from Chignahuapan, Puebla, Mexico Reveals the Presence of Particular Sulfur-Oxidizing Bacterial and Viral Communities. <i>Microorganisms</i> , 2020 , 8,	4.9	2
17	sp. Strain SGH1, a Bacterioruberin-Rich, Perchlorate-Tolerant Halophilic Archaeon Isolated From Halite Microbial Communities, Atacama Desert, Chile. <i>Frontiers in Microbiology</i> , 2020 , 11, 324	5.7	10
16	Stress Reshapes the Physiological Response of Halophile Fungi to Salinity. <i>Cells</i> , 2020 , 9,	7.9	15
15	Transcriptome during the Infection Process of the Bryophyte and Angiosperms. <i>Journal of Fungi (Basel, Switzerland)</i> , 2020 , 7,	5.6	4
14	Exogenous Nitric Oxide Delays Plant Regeneration from Protoplast and Protonema Development in. <i>Plants</i> , 2020 , 9,	4.5	1
13	Haloadaptative Responses of to Extreme Water Deprivation: Morphology, Compatible Solutes, and Oxidative Stress at NaCl Saturation. <i>Journal of Fungi (Basel, Switzerland)</i> , 2020 , 6,	5.6	5
12	Aromatic Hydrocarbon Removal by Novel Extremotolerant and Spp. from an Oil Polluted Site in Mexico. <i>Journal of Fungi (Basel, Switzerland)</i> , 2020 , 6,	5.6	4
11	First demonstration that ascomycetous halophilic fungi (<i>Aspergillus sydowii</i> and <i>Aspergillus destruens</i>) are useful in xenobiotic mycoremediation under high salinity conditions. <i>Bioresource Technology</i> , 2019 , 279, 287-296	11	33

10	Extremophile deep-sea viral communities from hydrothermal vents: Structural and functional analysis. <i>Marine Genomics</i> , 2019 , 46, 16-28	1.9	15
9	Intermediate-Salinity Systems at High Altitudes in the Peruvian Andes Unveil a High Diversity and Abundance of Bacteria and Viruses. <i>Genes</i> , 2019 , 10,	4.2	3
8	Metagenomics of Atacama Lithobiontic Extremophile Life Unveils Highlights on Fungal Communities, Biogeochemical Cycles and Carbohydrate-Active Enzymes. <i>Microorganisms</i> , 2019 , 7,	4.9	13
7	Schizophyllum commune: An unexploited source for lignocellulose degrading enzymes. <i>MicrobiologyOpen</i> , 2018 , 7, e00637	3.4	6
6	Simple screening protocol for identification of potential mycoremediation tools for the elimination of polycyclic aromatic hydrocarbons and phenols from hyperalkalophile industrial effluents. <i>Journal of Environmental Management</i> , 2017 , 198, 1-11	7.9	33
5	From lignocellulosic metagenomes to lignocellulolytic genes: trends, challenges and future prospects. <i>Biofuels, Bioproducts and Biorefining</i> , 2016 , 10, 864-882	5.3	30
4	Xenobiotic Compounds Degradation by Heterologous Expression of a Trametes sanguineus Laccase in Trichoderma atroviride. <i>PLoS ONE</i> , 2016 , 11, e0147997	3.7	40
3	Identification of a novel carbohydrate esterase from Bjerkandera adusta: structural and function predictions through bioinformatics analysis and molecular modeling. <i>Proteins: Structure, Function and Bioinformatics</i> , 2015 , 83, 533-46	4.2	6
2	A novel expansin protein from the white-rot fungus Schizophyllum commune. <i>PLoS ONE</i> , 2015 , 10, e0122296	3.7	27
1	Characterization of lignocellulolytic activities from a moderate halophile strain of Aspergillus caesiellus isolated from a sugarcane bagasse fermentation. <i>PLoS ONE</i> , 2014 , 9, e105893	3.7	22