

Tania Aires

List of Publications by Year in descending order

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Version: 2024-02-01

18
papers

595
citations

567281

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940533

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826
citing authors

#	ARTICLE	IF	CITATIONS
1	Host and Environmental Specificity in Bacterial Communities Associated to Two Highly Invasive Marine Species (Genus <i>Asparagopsis</i>). <i>Frontiers in Microbiology</i> , 2016, 7, 559.	3.5	72
2	Invasion Is a Community Affair: Clandestine Followers in the Bacterial Community Associated to Green Algae, <i>Caulerpa racemosa</i> , Track the Invasion Source. <i>PLoS ONE</i> , 2013, 8, e68429.	2.5	63
3	Glacial vicariance drives phylogeographic diversification in the amphi-boreal kelp <i>Saccharina latissima</i> . <i>Scientific Reports</i> , 2018, 8, 1112.	3.3	61
4	Summer shifts of bacterial communities associated with the invasive brown seaweed <i>Sargassum muticum</i> are location and tissue dependent. <i>PLoS ONE</i> , 2018, 13, e0206734.	2.5	57
5	Hologenome theory supported by cooccurrence networks of species-specific bacterial communities in siphonous algae (<i>Caulerpa</i>). <i>FEMS Microbiology Ecology</i> , 2015, 91, fiv067.	2.7	55
6	Species Specificity of Bacteria Associated to the Brown Seaweeds <i>Lobophora</i> (Dictyotales). <i>Frontiers in Microbiology</i> , 2016, 7, 316.	3.5	53
7	Evolutionary history of the seagrass genus <i>Posidonia</i> . <i>Marine Ecology - Progress Series</i> , 2011, 421, 117-130.	1.9	40
8	Expressed sequence tags from heat-shocked seagrass <i>Zostera noltii</i> (Hornemann) from its southern distribution range. <i>Marine Genomics</i> , 2011, 4, 181-188.	1.1	29
9	Host Differentiation and Compartmentalization of Microbial Communities in the Azooxanthellate Cupcorals <i>Tubastrea coccinea</i> and <i>Rhizopsammia goesi</i> in the Caribbean. <i>Frontiers in Marine Science</i> , 2018, 5, .	2.5	25
10	Seaweed Loads Cause Stronger Bacterial Community Shifts in Coastal Lagoon Sediments Than Nutrient Loads. <i>Frontiers in Microbiology</i> , 2018, 9, 3283.	3.5	25
11	Lyme borreliosis spirochetes in questing ticks from mainland Portugal. <i>International Journal of Medical Microbiology Supplements</i> , 2004, 293, 109-116.	0.4	23
12	The interaction between the proliferating macroalga <i>Asparagopsis taxiformis</i> and the coral <i>Astroides calycularis</i> induces changes in microbiome and metabolomic fingerprints. <i>Scientific Reports</i> , 2017, 7, 42625.	3.3	23
13	Characterization of 12 polymorphic microsatellite markers in the sugar kelp <i>Saccharina latissima</i> . <i>Journal of Applied Phycology</i> , 2016, 28, 3071-3074.	2.8	22
14	SELECTIVE ELIMINATION OF CHLOROPLASTIDIAL DNA FOR METAGENOMICS OF BACTERIA ASSOCIATED WITH THE GREEN ALGA <i>CAULERPA TAXIFOLIA</i> (BRYOPSIDOPHYCEAE). <i>Journal of Phycology</i> , 2012, 48, 483-490.	2.3	19
15	Acidification increases abundances of <i>Vibrionales</i> and <i>Planctomycetia</i> associated to a seaweed-grazer system: potential consequences for disease and prey digestion efficiency. <i>PeerJ</i> , 2018, 6, e4377.	2.0	16
16	Characterization and Comparison of Bacterial Communities of an Invasive and Two Native Caribbean Seagrass Species Sheds Light on the Possible Influence of the Microbiome on Invasive Mechanisms. <i>Frontiers in Microbiology</i> , 2021, 12, 653998.	3.5	10
17	Unraveling seaweeds bacteriomes. , 2018, , 95-113.		2
18	Microbial Surface Biofilm Responds to the Growth-Reproduction-Senescence Cycle of the Dominant Coral Reef Macroalgae <i>Sargassum</i> spp.. <i>Life</i> , 2021, 11, 1199.	2.4	0