## **Tania Aires**

List of Publications by Year in descending order

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		567281 940533	
18	595	15	16
papers	citations	h-index	g-index
18	18	18	826
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	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. Frontiers in Microbiology, 2021, 12, 653998.	3.5	10
2	Microbial Surface Biofilm Responds to the Growth-Reproduction-Senescence Cycle of the Dominant Coral Reef Macroalgae Sargassum spp Life, 2021, 11, 1199.	2.4	0
3	Glacial vicariance drives phylogeographic diversification in the amphi-boreal kelp Saccharina latissima. Scientific Reports, 2018, 8, 1112.	3.3	61
4	Host Differentiation and Compartmentalization of Microbial Communities in the Azooxanthellate Cupcorals Tubastrea coccinea and Rhizopsammia goesi in the Caribbean. Frontiers in Marine Science, 2018, 5, .	2.5	25
5	Summer shifts of bacterial communities associated with the invasive brown seaweed Sargassum muticum are location and tissue dependent. PLoS ONE, 2018, 13, e0206734.	2.5	57
6	Seaweed Loads Cause Stronger Bacterial Community Shifts in Coastal Lagoon Sediments Than Nutrient Loads. Frontiers in Microbiology, 2018, 9, 3283.	3.5	25
7	Unraveling seaweeds bacteriomes. , 2018, , 95-113.		2
8	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. PeerJ, 2018, 6, e4377.	2.0	16
9	The interaction between the proliferating macroalga Asparagopsis taxiformis and the coral Astroides calycularis induces changes in microbiome and metabolomic fingerprints. Scientific Reports, 2017, 7, 42625.	3.3	23
10	Species Specificity of Bacteria Associated to the Brown Seaweeds Lobophora (Dictyotales,) Tj ETQq0 0 0 rgBT /O Frontiers in Microbiology, 2016, 7, 316.	verlock 10 3.5	O Tf 50 387 Tc 53
11	Host and Environmental Specificity in Bacterial Communities Associated to Two Highly Invasive Marine Species (Genus Asparagopsis). Frontiers in Microbiology, 2016, 7, 559.	3.5	72
12	Characterization of 12 polymorphic microsatellite markers in the sugar kelp Saccharina latissima. Journal of Applied Phycology, 2016, 28, 3071-3074.	2.8	22
13	Hologenome theory supported by cooccurrence networks of species-specific bacterial communities in siphonous algae ( <i>Caulerpa</i> ). FEMS Microbiology Ecology, 2015, 91, fiv067.	2.7	55
14	Invasion Is a Community Affair: Clandestine Followers in the Bacterial Community Associated to Green Algae, Caulerpa racemosa, Track the Invasion Source. PLoS ONE, 2013, 8, e68429.	2.5	63
15	SELECTIVE ELIMINATION OF CHLOROPLASTIDIAL DNA FOR METAGENOMICS OF BACTERIA ASSOCIATED WITH THE GREEN ALGA <i>CAULERPA TAXIFOLIA</i> (BRYOPSIDOPHYCEAE) Sup>1 Journal of Phycology, 2012, 48, 483-490.	2.3	19
16	Expressed sequence tags from heat-shocked seagrass Zostera noltii (Hornemann) from its southern distribution range. Marine Genomics, 2011, 4, 181-188.	1,1	29
17	Evolutionary history of the seagrass genus Posidonia. Marine Ecology - Progress Series, 2011, 421, 117-130.	1.9	40
18	Lyme borreliosis spirochetes in questing ticks from mainland Portugal. International Journal of Medical Microbiology Supplements, 2004, 293, 109-116.	0.4	23