Daniel Cerqueda-GarcÃ-a

List of Publications by Year in descending order

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Version: 2024-02-01

687220 839398 31 440 13 18 g-index citations h-index papers 37 37 37 660 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	Molecular Docking and Dynamics Simulation of Protein \hat{I}^2 -Tubulin and Antifungal Cyclic Lipopeptides. Molecules, 2019, 24, 3387.	1.7	34
2	Exploring Biogeochemistry and Microbial Diversity of Extant Microbialites in Mexico and Cuba. Frontiers in Microbiology, 2018, 9, 510.	1.5	29
3	Temperature susceptibility of a mesophilic anaerobic membrane bioreactor treating saline phenol-containing wastewater. Chemosphere, 2018, 213, 92-102.	4.2	27
4	Assessment of the bacterial community structure in shallow and deep sediments of the Perdido Fold Belt region in the Gulf of Mexico. PeerJ, 2018, 6, e5583.	0.9	26
5	Disturbance in human gut microbiota networks by parasites and its implications in the incidence of depression. Scientific Reports, 2020, 10, 3680.	1.6	22
6	Toxicity evaluation and microbiota response of the lined sole Achirus lineatus (Chordata: Achiridae) exposed to the light petroleum water-accommodated fraction (WAF). Journal of Toxicology and Environmental Health - Part A: Current Issues, 2020, 83, 313-329.	1.1	21
7	Bacterial succession and coâ€occurrence patterns of an enriched marine microbial community during light crude oil degradation in a batch reactor. Journal of Applied Microbiology, 2019, 127, 495-507.	1.4	20
8	Microbial distribution and turnover in Antarctic microbial mats highlight the relevance of heterotrophic bacteria in low-nutrient environments. FEMS Microbiology Ecology, 2018, 94, .	1.3	19
9	Changes in the Bacterioplankton Community Structure from Southern Gulf of Mexico During a Simulated Crude Oil Spill at Mesocosm Scale. Microorganisms, 2019, 7, 441.	1.6	18
10	A succession of marine bacterial communities in batch reactor experiments during the degradation of five different petroleum types. Marine Pollution Bulletin, 2020, 150, 110775.	2.3	17
11	Effects of chronic exposure to water accommodated fraction (WAF) of light crude oil on gut microbiota composition of the lined sole (Achirus lineatus). Marine Environmental Research, 2020, 161, 105116.	1.1	17
12	Local dynamics of a white syndrome outbreak and changes in the microbial community associated with colonies of the scleractinian brain coral <i>Pseudodiploria strigosa</i> PeerJ, 2021, 9, e10695.	0.9	17
13	Fecal microbiota of different reproductive stages of the central population of the lesser-long nosed bat, Leptonycteris yerbabuenae. PLoS ONE, 2019, 14, e0219982.	1.1	15
14	Geographical separation and physiology drive differentiation of microbial communities of two discrete populations of the bat <i>Leptonycteris yerbabuenae</i> i>. MicrobiologyOpen, 2020, 9, 1113-1127.	1.2	15
15	Metabolic analysis of <i>Chlorobium chlorochromatii</i> CaD3 reveals clues of the symbiosis in $\hat{a} \in \tilde{a}$ i>Chlorochromatium aggregatum $\hat{a} \in \tilde{a}$. ISME Journal, 2014, 8, 991-998.	4.4	13
16	Microbiota composition of the dorsal patch of reproductive male Leptonycteris yerbabuenae. PLoS ONE, 2019, 14, e0226239.	1.1	13
17	Gut Microbiome in Children from Indigenous and Urban Communities in México: Different Subsistence Models, Different Microbiomes. Microorganisms, 2020, 8, 1592.	1.6	13
18	Metabolic potential of microbial mats and microbialites: Autotrophic capabilities described by an <i>in silico</i> stoichiometric approach from shared genomic resources. Journal of Bioinformatics and Computational Biology, 2016, 14, 1650020.	0.3	11

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19	Anaerobic Conversion of Saline Phenol-Containing Wastewater Under Thermophilic Conditions in a Membrane Bioreactor. Frontiers in Bioengineering and Biotechnology, 2020, 8, 565311.	2.0	11
20	Alterations in the Gut Microbiota of Zebrafish (Danio rerio) in Response to Water-Soluble Crude Oil Components and Its Mixture With a Chemical Dispersant. Frontiers in Public Health, 2020, 8, 584953.	1.3	11
21	Towards an understanding of the role of intrinsic protein disorder on plant adaptation to environmental challenges. Cell Stress and Chaperones, 2021, 26, 141-150.	1.2	11
22	Microbial composition of biofilms associated with lithifying rubble of Acropora palmatabranches. FEMS Microbiology Ecology, 2016, 92, fiv162.	1.3	10
23	Enhancing Phenol Conversion Rates in Saline Anaerobic Membrane Bioreactor Using Acetate and Butyrate as Additional Carbon and Energy Sources. Frontiers in Microbiology, 2020, 11, 604173.	1.5	10
24	Degradation of p-cresol, resorcinol, and phenol in anaerobic membrane bioreactors under saline conditions. Chemical Engineering Journal, 2022, 430, 132672.	6.6	9
25	Community structure and distribution of benthic Bacteria and Archaea in a stratified coastal lagoon in the Southern Gulf of Mexico. Estuarine, Coastal and Shelf Science, 2019, 230, 106433.	0.9	7
26	Assessing the Diversity of Benthic Sulfate-Reducing Microorganisms in Northwestern Gulf of Mexico by Illumina Sequencing of dsrB Gene. Microbial Ecology, 2021, 81, 908-921.	1.4	6
27	Alterations in the gut-associated microbiota of juvenile Caribbean spiny lobsters Panulirus argus (Latreille, 1804) infected with PaV1. Journal of Invertebrate Pathology, 2020, 176, 107457.	1.5	5
28	Effects of a Light Crude Oil Spill on a Tropical Coastal Phytoplankton Community. Bulletin of Environmental Contamination and Toxicology, 2022, 108, 55-63.	1.3	5
29	Assessing the Effect of Chemical Dispersant Nokomis 3-F4 on the Degradation of a Heavy Crude Oil in Water by a Marine Microbial Consortium. Bulletin of Environmental Contamination and Toxicology, 2021, , 1.	1.3	1
30	First record of entomopathogenic nematodes from Yucat \tilde{A}_i n State, M \tilde{A} $\hat{\mathbb{Q}}$ xico and their infectivity capacity against <i>Aedes aegypti</i> . PeerJ, 2021, 9, e11633.	0.9	1
31	Alterations in the sap-associated microbiota of Carica papaya in response to drought stress. Symbiosis, 2020, 81, 93-100.	1.2	O