Jenan J Kharbush

List of Publications by Year in descending order

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IENAN I KHADRUSH

#	Article	IF	CITATIONS
1	Sharing and community curation of mass spectrometry data with Global Natural Products Social Molecular Networking. Nature Biotechnology, 2016, 34, 828-837.	9.4	2,802
2	Particulate Organic Carbon Deconstructed: Molecular and Chemical Composition of Particulate Organic Carbon in the Ocean. Frontiers in Marine Science, 2020, 7, .	1.2	72
3	Composite Bacterial Hopanoids and Their Microbial Producers across Oxygen Gradients in the Water Column of the California Current. Applied and Environmental Microbiology, 2013, 79, 7491-7501.	1.4	29
4	Intact polar diacylglycerol biomarker lipids isolated from suspended particulate organic matter accumulating in an ultraoligotrophic water column. Organic Geochemistry, 2016, 100, 29-41.	0.9	17
5	Vitamin B ₁₂ -dependent biosynthesis ties amplified 2-methylhopanoid production during oceanic anoxic events to nitrification. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 32996-33004.	3.3	13
6	Chlorophyll nitrogen isotope values track shifts between cyanobacteria and eukaryotic algae in a natural phytoplankton community in Lake Erie. Organic Geochemistry, 2019, 128, 71-77.	0.9	9
7	Distribution and Abundance of Hopanoid Producers in Low-Oxygen Environments of the Eastern Pacific Ocean. Microbial Ecology, 2016, 71, 401-408.	1.4	8
8	Hopanoid-producing bacteria in the Red Sea include the major marine nitrite oxidizers. FEMS Microbiology Ecology, 2018, 94, .	1.3	8
9	Linking diatom-diazotroph symbioses to nitrogen cycle perturbations and deep-water anoxia: Insights from Mediterranean sapropel events. Earth and Planetary Science Letters, 2021, 571, 117110.	1.8	8
10	Marine and terrestrial nitrifying bacteria are sources of diverse bacteriohopanepolyols. Geobiology, 2022, 20, 399-420.	1.1	8
11	Composite Bacterial Hopanoids and Their Microbial Producers across Oxygen Gradients in the Water Column of the California Current. Applied and Environmental Microbiology, 2014, 80, 3283-3283.	1.4	0