

Anchittha Satjarak

List of Publications by Year in descending order

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#	ARTICLE	IF	CITATIONS
1	Plastid Genome of <i>Equisetum xylochaetum</i> from the Atacama Desert, Chile and the Relationships of <i>Equisetum</i> Based on Frequently Used Plastid Genes and Network Analysis. <i>Plants</i> , 2022, 11, 1001.	3.5	0
2	Microbiome and related structural features of Earth's most archaic plant indicate early plant symbiosis attributes. <i>Scientific Reports</i> , 2022, 12, 6423.	3.3	3
3	Nitrogen fixation and other biogeochemically important features of Atacama Desert giant horsetail plant microbiomes inferred from metagenomic contig analysis. <i>Annals of Botany</i> , 2022, , .	2.9	2
4	Shotgun metagenomics and microscopy indicate diverse cyanophytes, other bacteria, and microeukaryotes in the epimicrobiota of a northern Chilean wetland <i>Nostoc</i> (Cyanobacteria). <i>Journal of Phycology</i> , 2021, 57, 39-50.	2.3	3
5	Bacteria Associated with <i>Echinodorus cordifolius</i> and <i>Lepironia articulata</i> Enhance Nitrogen and Phosphorus Removal from Wastewater. <i>Bulletin of Environmental Contamination and Toxicology</i> , 2021, 106, 377-384.	2.7	3
6	The symbiosis between <i>Philidris</i> ants and the ant-plant <i>Dischidia major</i> includes fungal and algal associates. <i>Symbiosis</i> , 2021, 83, 305-315.	2.3	3
7	Hercide Atrazine Alters the Microbiota of the Filamentous Green Alga <i>Cladophora</i> sp. Cultured from Thailand. <i>Sains Malaysiana</i> , 2021, 50, 1255-1265.	0.5	1
8	Microscopic and Metagenomic Evidence for Eukaryotic Microorganisms Associated with Atacama Desert Populations of Giant <i>Equisetum</i> . <i>American Fern Journal</i> , 2021, 111, .	0.3	3
9	A biorefinery approach for high value-added bioproduct (astaxanthin) from alga <i>Haematococcus</i> sp. and residue pyrolysis for biochar synthesis and metallic iron production from hematite (Fe ₂ O ₃). <i>Fuel</i> , 2021, 304, 121150.	6.4	9
10	Spatial Variation of <i>Cladophora</i> Epiphytes in the Nan River, Thailand. <i>Plants</i> , 2021, 10, 2266.	3.5	2
11	Molecular markers obtained from draft genomic sequence data characterise an isolate of <i>Oedogonium</i> (Chlorophyceae) used for biomass applications. <i>Phycologia</i> , 2020, 59, 340-345.	1.4	1
12	Evolution of organellar genes of chlorophyte algae: Relevance to phylogenetic inference. <i>PLoS ONE</i> , 2019, 14, e0216608.	2.5	4
13	Comparative <i>scn</i> DNA <i>scn</i> sequence analyses of <i>Pyramimonas parkeae</i> (Prasinophyceae) chloroplast genomes. <i>Journal of Phycology</i> , 2017, 53, 415-424.	2.3	9
14	Complete mitochondrial genomes of prasinophyte algae <i>Pyramimonas parkeae</i> and <i>Cymbomonas tetramitiformis</i> . <i>Journal of Phycology</i> , 2017, 53, 601-615.	2.3	10
15	Genome-wide analysis of carbohydrate-active enzymes in <i>Pyramimonas parkeae</i> (Prasinophyceae). <i>Journal of Phycology</i> , 2017, 53, 1072-1086.	2.3	7
16	Complete Chloroplast Genome Sequence of Phagomixotrophic Green Alga <i>Cymbomonas tetramitiformis</i> . <i>Genome Announcements</i> , 2016, 4, .	0.8	8