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List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	The Phaeodactylum genome reveals the evolutionary history of diatom genomes. Nature, 2008, 456, 239-244.	27.8	1,458
2	Molecular toolbox for studying diatom biology in Phaeodactylum tricornutum. Gene, 2007, 406, 23-35.	2.2	293
3	An atypical member of the light-harvesting complex stress-related protein family modulates diatom responses to light. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 18214-18219.	7.1	258
4	Diatom PtCPF1 is a new cryptochrome/photolyase family member with DNA repair and transcription regulation activity. EMBO Reports, 2009, 10, 655-661.	4.5	168
5	Evolutionary Origins and Functions of the Carotenoid Biosynthetic Pathway in Marine Diatoms. PLoS ONE, 2008, 3, e2896.	2.5	134
6	Genome-wide diel growth state transitions in the diatom <i>Thalassiosira pseudonana</i> . Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 7518-7523.	7.1	132
7	Sulfonate-based networks between eukaryotic phytoplankton and heterotrophic bacteria in the surface ocean. Nature Microbiology, 2019, 4, 1706-1715.	13.3	120
8	Nutrient Limitation is the Main Regulatory Factor for Carotenoid Accumulation and for Psy and Pds Steady State Transcript Levels in Dunaliella salina (Chlorophyta) Exposed to High Light and Salt Stress. Marine Biotechnology, 2008, 10, 602-11.	2.4	110
9	IDENTIFICATION AND COMPARATIVE GENOMIC ANALYSIS OF SIGNALING AND REGULATORY COMPONENTS IN THE DIATOMTHALASSIOSIRA PSEUDONANA. Journal of Phycology, 2007, 43, 585-604.	2.3	87
10	Isolation and characterization of a stress-inducible Dunaliella salina Lcy-β gene encoding a functional lycopene β-cyclase. Applied Microbiology and Biotechnology, 2008, 79, 819-28.	3.6	65
11	The dynamic trophic architecture of open-ocean protist communities revealed through machine-guided metatranscriptomics. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, .	7.1	27
12	Diel transcriptional oscillations of light-sensitive regulatory elements in open-ocean eukaryotic plankton communities. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	24
13	Complex marine microbial communities partition metabolism of scarce resources over the diel cycle. Nature Ecology and Evolution, 2022, 6, 218-229.	7.8	21
14	Fe limitation decreases transcriptional regulation over the diel cycle in the model diatom Thalassiosira pseudonana. PLoS ONE, 2019, 14, e0222325.	2.5	14
15	Diel-Regulated Transcriptional Cascades of Microbial Eukaryotes in the North Pacific Subtropical Gyre. Frontiers in Microbiology, 2021, 12, 682651.	3.5	6