

Hervé Moreau

List of Publications by Year in descending order

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43
papers

5,025
citations

172457

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254184

43
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46
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docs citations

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times ranked

4907
citing authors

#	ARTICLE	IF	CITATIONS
1	<i>Mantoniella beaufortii</i> and <i>Mantoniella baffinensis</i> sp. nov. (Mamiellales,) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 507 Phycology, 2020, 56, 37-51.	2.3	14
2	Virus-host coexistence in phytoplankton through the genomic lens. Science Advances, 2020, 6, eaay2587.	10.3	30
3	Simplified Transformation of <i>Ostreococcus tauri</i> Using Polyethylene Glycol. Genes, 2019, 10, 399.	2.4	14
4	Prasinovirus Attack of <i>Ostreococcus</i> Is Furtive by Day but Savage by Night. Journal of Virology, 2018, 92, .	3.4	42
5	Rapidity of Genomic Adaptations to Prasinovirus Infection in a Marine Microalga. Viruses, 2018, 10, 441.	3.3	10
6	Genome Analyses of the Microalga <i>Picochlorum</i> Provide Insights into the Evolution of Thermotolerance in the Green Lineage. Genome Biology and Evolution, 2018, 10, 2347-2365.	2.5	36
7	Host-derived viral transporter protein for nitrogen uptake in infected marine phytoplankton. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E7489-E7498.	7.1	74
8	Population genomics of picophytoplankton unveils novel chromosome hypervariability. Science Advances, 2017, 3, e1700239.	10.3	73
9	A Viral Immunity Chromosome in the Marine Picoeukaryote, <i>Ostreococcus tauri</i> . PLoS Pathogens, 2016, 12, e1005965.	4.7	38
10	Diversity of Viruses Infecting the Green Microalga <i>Ostreococcus lucimarinus</i> . Journal of Virology, 2015, 89, 5812-5821.	3.4	35
11	Prasinovirus distribution in the Northwest Mediterranean Sea is affected by the environment and particularly by phosphate availability. Virology, 2014, 466-467, 146-157.	2.4	17
12	An improved genome of the model marine alga <i>Ostreococcus tauri</i> unfolds by assessing Illumina de novo assemblies. BMC Genomics, 2014, 15, 1103.	2.8	90
13	Morphology, Genome Plasticity, and Phylogeny in the Genus <i>Ostreococcus</i> Reveal a Cryptic Species, <i>O. mediterraneus</i> sp. nov. (Mamiellales, Mamiellophyceae). Protist, 2013, 164, 643-659.	1.5	48
14	Evolution of Codon Usage in the Smallest Photosynthetic Eukaryotes and Their Giant Viruses. Genome Biology and Evolution, 2013, 5, 848-859.	2.5	24
15	picoâ€œscp>PLAZA</scp>, a genome database of microbial photosynthetic eukaryotes. Environmental Microbiology, 2013, 15, 2147-2153.	3.8	87
16	Exploring nucleo-cytoplasmic large DNA viruses in Tara Oceans microbial metagenomes. ISME Journal, 2013, 7, 1678-1695.	9.8	185
17	Strategies and mechanisms of resistance to viruses in photosynthetic aquatic microorganisms. Advances in Oceanography and Limnology, 2012, 3, 1-15.	0.6	5
18	Gene functionalities and genome structure in <i>Bathycoccus prasinus</i> reflect cellular specializations at the base of the green lineage. Genome Biology, 2012, 13, R74.	9.6	143

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19	Genomics of Algal Host-Virus Interactions. <i>Advances in Botanical Research</i> , 2012, , 343-381.	1.1	15
20	Phylogeny and Molecular Evolution of the Green Algae. <i>Critical Reviews in Plant Sciences</i> , 2012, 31, 1-46.	5.7	723
21	Genome diversity in the smallest marine photosynthetic eukaryotes. <i>Research in Microbiology</i> , 2011, 162, 570-577.	2.1	33
22	Acquisition and maintenance of resistance to viruses in eukaryotic phytoplankton populations. <i>Environmental Microbiology</i> , 2011, 13, 1412-1420.	3.8	70
23	How and Why DNA Barcodes Underestimate the Diversity of Microbial Eukaryotes. <i>PLoS ONE</i> , 2011, 6, e16342.	2.5	62
24	Marine Prasinovirus Genomes Show Low Evolutionary Divergence and Acquisition of Protein Metabolism Genes by Horizontal Gene Transfer. <i>Journal of Virology</i> , 2010, 84, 12555-12563.	3.4	87
25	Abundance, spatial distribution and genetic diversity of <i>Ostreococcus tauri</i> viruses in two different environments. <i>Environmental Microbiology Reports</i> , 2010, 2, 313-321.	2.4	32
26	Cryptic Sex in the Smallest Eukaryotic Marine Green Alga. <i>Molecular Biology and Evolution</i> , 2010, 27, 47-54.	8.9	81
27	Unravelling cis-Regulatory Elements in the Genome of the Smallest Photosynthetic Eukaryote: Phylogenetic Footprinting in <i>Ostreococcus</i> . <i>Journal of Molecular Evolution</i> , 2009, 69, 249-259.	1.8	10
28	Green Evolution and Dynamic Adaptations Revealed by Genomes of the Marine Picoeukaryotes <i>Micromonas</i> . <i>Science</i> , 2009, 324, 268-272.	12.6	591
29	Phylogenetic analysis of new Prasinoviruses (<i>Phycodnaviridae</i>) that infect the green unicellular algae <i>Ostreococcus</i> , <i>Bathycoccus</i> and <i>Micromonas</i> . <i>Environmental Microbiology Reports</i> , 2009, 1, 114-123.	2.4	35
30	Picoeukaryotic sequences in the Sargasso Sea metagenome. <i>Genome Biology</i> , 2008, 9, R5.	9.6	34
31	Clues about the Genetic Basis of Adaptation Emerge from Comparing the Proteomes of Two <i>Ostreococcus</i> Ecotypes (Chlorophyta, Prasinophyceae). <i>Molecular Biology and Evolution</i> , 2008, 25, 2293-2300.	8.9	39
32	An original adaptation of photosynthesis in the marine green alga <i>Ostreococcus</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 7881-7886.	7.1	154
33	Life-Cycle and Genome of OtV5, a Large DNA Virus of the Pelagic Marine Unicellular Green Alga <i>Ostreococcus tauri</i> . <i>PLoS ONE</i> , 2008, 3, e2250.	2.5	107
34	The tiny eukaryote <i>Ostreococcus</i> provides genomic insights into the paradox of plankton speciation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 7705-7710.	7.1	563
35	The Complete Chloroplast and Mitochondrial DNA Sequence of <i>Ostreococcus tauri</i> : Organelle Genomes of the Smallest Eukaryote Are Examples of Compaction. <i>Molecular Biology and Evolution</i> , 2007, 24, 956-968.	8.9	106
36	Screening the Sargasso Sea metagenome for data to investigate genome evolution in <i>Ostreococcus</i> (Prasinophyceae, Chlorophyta). <i>Gene</i> , 2007, 406, 184-190.	2.2	28

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37	Genome analysis of the smallest free-living eukaryote <i>Ostreococcus tauri</i> unveils many unique features. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 11647-11652.	7.1	809
38	Natural Synchronisation for the Study of Cell Division in the Green Unicellular Alga <i>Ostreococcus tauri</i> . <i>Plant Molecular Biology</i> , 2006, 60, 277-292.	3.9	36
39	Ecotype diversity in the marine picoeukaryote <i>Ostreococcus</i> (Chlorophyta, Prasinophyceae). <i>Environmental Microbiology</i> , 2005, 7, 853-859.	3.8	185
40	Genome-Wide Analysis of Core Cell Cycle Genes in the Unicellular Green Alga <i>Ostreococcus tauri</i> . <i>Molecular Biology and Evolution</i> , 2005, 22, 589-597.	8.9	64
41	New Insights into the Nature and Phylogeny of Prasinophyte Antenna Proteins: <i>Ostreococcus tauri</i> , a Case Study. <i>Molecular Biology and Evolution</i> , 2005, 22, 2217-2230.	8.9	69
42	Starch Division and Partitioning. A Mechanism for Granule Propagation and Maintenance in the Picophytoplanktonic Green Alga <i>Ostreococcus tauri</i> . <i>Plant Physiology</i> , 2004, 136, 3333-3340.	4.8	80
43	DNA LIBRARIES FOR SEQUENCING THE GENOME OF <i>Ostreococcus tauri</i> (Chlorophyta.) <i>Tj ETQq1</i> 1 0.784314 rgBT /Overlock 10 1150-1156.	2.3	42