Laure Guillou

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7675647/publications.pdf

Version: 2024-02-01

72 papers

8,424 citations

38 h-index 71 g-index

77 all docs

77 docs citations

77 times ranked 7031 citing authors

#	Article	IF	CITATIONS
1	The Protist Ribosomal Reference database (PR2): a catalog of unicellular eukaryote Small Sub-Unit rRNA sequences with curated taxonomy. Nucleic Acids Research, 2012, 41, D597-D604.	6.5	1,463
2	Revisions to the Classification, Nomenclature, and Diversity of Eukaryotes. Journal of Eukaryotic Microbiology, 2019, 66, 4-119.	0.8	904
3	Widespread occurrence and genetic diversity of marine parasitoids belonging to <i>Syndiniales</i> (<i>Alveolata</i>). Environmental Microbiology, 2008, 10, 3349-3365.	1.8	511
4	Marine protist diversity in <scp>E</scp> uropean coastal waters and sediments as revealed by highâ€throughput sequencing. Environmental Microbiology, 2015, 17, 4035-4049.	1.8	384
5	Control of Toxic Marine Dinoflagellate Blooms by Serial Parasitic Killers. Science, 2008, 322, 1254-1257.	6.0	322
6	Phylogenetic and Ecological Analysis of Novel Marine Stramenopiles. Applied and Environmental Microbiology, 2004, 70, 3528-3534.	1.4	321
7	Diversity of Picoplanktonic Prasinophytes Assessed by Direct Nuclear SSU rDNA Sequencing of Environmental Samples and Novel Isolates Retrieved from Oceanic and Coastal Marine Ecosystems. Protist, 2004, 155, 193-214.	0.6	235
8	BOLIDOMONAS: A NEW GENUS WITH TWO SPECIES BELONGING TO A NEW ALGAL CLASS, THE BOLIDOPHYCEAE (HETEROKONTA). Journal of Phycology, 1999, 35, 368-381.	1.0	225
9	Abundance and diversity of prymnesiophytes in the picoplankton col_{4} munity from the equatorial Pacific Ocean inferred from 18S rDNA sequences. Limnology and Oceanography, 2000, 45, 98-109.	1.6	208
10	Eukaryotic Richness in the Abyss: Insights from Pyrotag Sequencing. PLoS ONE, 2011, 6, e18169.	1.1	207
11	Picoeukaryotic diversity in an oligotrophic coastal site studied by molecular and culturing approaches. FEMS Microbiology Ecology, 2004, 50, 231-243.	1.3	204
12	Phyto <scp>REF</scp> : a reference database of the plastidial 16S <scp>rRNA</scp> gene of photosynthetic eukaryotes with curated taxonomy. Molecular Ecology Resources, 2015, 15, 1435-1445.	2.2	198
13	Validation and Application of a PCR Primer Set to Quantify Fungal Communities in the Soil Environment by Real-Time Quantitative PCR. PLoS ONE, 2011, 6, e24166.	1.1	198
14	Ecotype diversity in the marine picoeukaryote Ostreococcus (Chlorophyta, Prasinophyceae). Environmental Microbiology, 2005, 7, 853-859.	1.8	185
15	Unveiling the Organisms behind Novel Eukaryotic Ribosomal DNA Sequences from the Ocean. Applied and Environmental Microbiology, 2002, 68, 4554-4558.	1.4	176
16	Oligonucleotide Probes for the Identification of Three Algal Groups by Dot Blot and Fluorescent Whole-Cell Hybridization. Journal of Eukaryotic Microbiology, 2000, 47, 76-84.	0.8	142
17	Phylogenetic analysis of the †Nannochloris-like' algae and diagnoses of Picochlorum oklahomensis gen. et sp. nov. (Trebouxiophyceae, Chlorophyta). Phycologia, 2004, 43, 641-652.	0.6	139
18	Rapid protein evolution, organellar reductions, and invasive intronic elements in the marine aerobic parasite dinoflagellate Amoebophrya spp. BMC Biology, 2021, 19, 1.	1.7	135

#	Article	IF	CITATIONS
19	Genetic Diversity and Molecular Detection of Three Toxic Dinoflagellate Genera and from French Coasts. Protist, 2002, 153, 223-238.	0.6	132
20	Wide genetic diversity of picoplanktonic green algae (Chloroplastida) in the Mediterranean Sea uncovered by a phylumâ€biased PCR approach. Environmental Microbiology, 2008, 10, 1804-1822.	1.8	112
21	Zoosporic parasites infecting marine diatoms – A black box that needs to be opened. Fungal Ecology, 2016, 19, 59-76.	0.7	109
22	Dinomyces arenysensis gen. et sp. nov. (Rhizophydiales, Dinomycetaceae fam. nov.), a Chytrid Infecting Marine Dinoflagellates. Protist, 2014, 165, 230-244.	0.6	102
23	EukRef: Phylogenetic curation of ribosomal RNA to enhance understanding of eukaryotic diversity and distribution. PLoS Biology, 2018, 16, e2005849.	2.6	101
24	PIGMENT SUITES AND TAXONOMIC GROUPS IN PRASINOPHYCEAE. Journal of Phycology, 2004, 40, 1149-1155.	1.0	99
25	IDENTIFICATION OF THE CLASS PRYMNESIOPHYCEAE AND THE GENUS PHAEOCYSTIS WITH RIBOSOMAL RNA-TARGETED NUCLEIC ACID PROBES DETECTED BY FLOW CYTOMETRY1. Journal of Phycology, 1996, 32, 858-868.	1.0	82
26	Comparative analysis between protist communities from the deepâ€sea pelagic ecosystem and specific deep hydrothermal habitats. Environmental Microbiology, 2010, 12, 2946-2964.	1.8	80
27	An aerobic eukaryotic parasite with functional mitochondria that likely lacks a mitochondrial genome. Science Advances, 2019, 5, eaav1110.	4.7	76
28	Diel variations in <i>Prochlorococcus</i> optical properties. Limnology and Oceanography, 2002, 47, 1637-1647.	1.6	75
29	Ecological impacts of parasitic chytrids, syndiniales and perkinsids on populations of marine photosynthetic dinoflagellates. Fungal Ecology, 2016, 19, 47-58.	0.7	7 3
30	PHYLOGENETIC ANALYSIS AND GENOME SIZE OF OSTREOCOCCUS TAURI (CHLOROPHYTA,) Tj ETQq0 0 0 rgBT /0	Overlock 1	0 ₇₂ 50 302
31	Diversity and Abundance of Bolidophyceae (Heterokonta) in Two Oceanic Regions. Applied and Environmental Microbiology, 1999, 65, 4528-4536.	1.4	72
32	The Roscoff Culture Collection (RCC): a collection dedicated to marine picoplankton. Nova Hedwigia, 2004, 79, 49-70.	0.2	71
33	Parvilucifera rostrata sp. nov. (Perkinsozoa), a Novel Parasitoid that Infects Planktonic Dinoflagellates. Protist, 2014, 165, 31-49.	0.6	69
34	Interplay Between the Parasite Amoebophrya sp. (Alveolata) and the Cyst Formation of the Red Tide Dinoflagellate Scrippsiella trochoidea. Protist, 2011, 162, 637-649.	0.6	64
35	General Patterns of Diversity in Major Marine Microeukaryote Lineages. PLoS ONE, 2013, 8, e57170.	1.1	54
36	Symbiomonas scintillans gen. et sp. nov. and Picophagus flagellatus gen. et sp. nov. (Heterokonta): Two New Heterotrophic Flagellates of Picoplanktonic Size. Protist, 1999, 150, 383-398.	0.6	53

#	Article	IF	Citations
37	Potential roles for recently discovered chytrid parasites in the dynamics of harmful algal blooms. Fungal Biology Reviews, 2015, 29, 20-33.	1.9	51
38	The Parasitic Dinoflagellates Blastodinium spp. Inhabiting the Gut of Marine, Planktonic Copepods: Morphology, Ecology, and Unrecognized Species Diversity. Frontiers in Microbiology, 2012, 3, 305.	1.5	45
39	Genomic Insights into Processes Driving the Infection of Alexandrium tamarense by the Parasitoid Amoebophrya sp. Eukaryotic Cell, 2014, 13, 1439-1449.	3.4	42
40	Molecular Diversity of the Syndinean Genus Euduboscquella Based on Single-Cell PCR Analysis. Applied and Environmental Microbiology, 2012, 78, 334-345.	1.4	40
41	Ultrastructure of Selenidium pendula, the Type Species of Archigregarines, and Phylogenetic Relations to Other Marine Apicomplexa. Protist, 2016, 167, 339-368.	0.6	40
42	Genetic diversity of Amoebophryidae (Syndiniales) during Alexandrium catenella/tamarense (Dinophyceae) blooms in the Thau lagoon (Mediterranean Sea, France). Research in Microbiology, 2011, 162, 959-968.	1.0	39
43	Protists Within Corals: The Hidden Diversity. Frontiers in Microbiology, 2018, 9, 2043.	1.5	39
44	Phylogenetic and morphological characterisation of the green algae infesting blue mussel <i>Mytilus edulis</i> in the North and South Atlantic oceans. Diseases of Aquatic Organisms, 2008, 81, 231-240.	0.5	39
45	Evidence for parasite-mediated selection during short-lasting toxic algal blooms. Proceedings of the Royal Society B: Biological Sciences, 2016, 283, 20161870.	1.2	38
46	A new view on the morphology and phylogeny of eugregarines suggested by the evidence from the gregarine <i>Ancora sagittata</i> (Leuckart, 1860) LabbA©, 1899 (Apicomplexa: Eugregarinida). PeerJ, 2017, 5, e3354.	0.9	29
47	Paralytic shellfish toxin content is related to genomic sxtA4 copy number in Alexandrium minutum strains. Frontiers in Microbiology, 2015, 6, 404.	1.5	28
48	Cryptic species in the parasitic Amoebophrya species complex revealed by a polyphasic approach. Scientific Reports, 2020, 10, 2531.	1.6	28
49	Evolutionary processes and cellular functions underlying divergence in <i>Alexandrium minutum</i> . Molecular Ecology, 2016, 25, 5129-5143.	2.0	25
50	Ecological and evolutionary significance of novel protist lineages. European Journal of Protistology, 2016, 55, 4-11.	0.5	25
51	Transcriptomic profiling of <i>Alexandrium fundyense</i> during physical interaction with or exposure to chemical signals from the parasite <i>Amoebophrya</i> . Molecular Ecology, 2016, 25, 1294-1307.	2.0	22
52	Does environmental heterogeneity explain temporal \hat{l}^2 diversity of small eukaryotic phytoplankton? Example from a tropical eutrophic coastal lagoon. Journal of Plankton Research, 2017, 39, 698-714.	0.8	21
53	CHARACTERIZATION OF THE PARMALES: MUCH MORE THAN THE RESOLUTION OF A TAXONOMIC ENIGMA. Journal of Phycology, 2011, 47, 2-4.	1.0	19
54	Comparative Time-Scale Gene Expression Analysis Highlights the Infection Processes of Two Amoebophrya Strains. Frontiers in Microbiology, 2018, 9, 2251.	1.5	19

#	Article	IF	CITATIONS
55	From the sxtA4 Gene to Saxitoxin Production: What Controls the Variability Among Alexandrium minutum and Alexandrium pacificum Strains?. Frontiers in Microbiology, 2021, 12, 613199.	1.5	19
56	Significance of Plankton Community Structure and Nutrient Availability for the Control of Dinoflagellate Blooms by Parasites: A Modeling Approach. PLoS ONE, 2015, 10, e0127623.	1.1	18
57	Fine structure and Molecular Phylogenetic Position of Two Marine Gregarines, Selenidium pygospionis sp. n. and S. pherusae sp. n., with Notes on the Phylogeny of Archigregarinida (Apicomplexa). Protist, 2018, 169, 826-852.	0.6	16
58	First Ultrastructural and Molecular Phylogenetic Evidence from the Blastogregarines, an Early Branching Lineage of Plesiomorphic Apicomplexa. Protist, 2018, 169, 697-726.	0.6	14
59	Identification to species level of live single microalgal cells from plankton samples with matrix-free laser/desorption ionization mass spectrometry. Metabolomics, 2020, 16, 28.	1.4	14
60	Photosymbiosis in Marine Pelagic Environments. , 2016, , 305-332.		13
61	Parasitic infections by Group <scp>II</scp> Syndiniales target selected dinoflagellate host populations within diverse protist assemblages in a model coastal pond. Environmental Microbiology, 2022, 24, 1818-1834.	1.8	13
62	16S rRNA gene-based molecular analysis of mat-forming and accompanying bacteria covering organically-enriched marine sediments underlying a salmon farm in Southern Chile (Calbuco Island). Gayana, 2010, 74, 125-1235.	0.0	10
63	Intracellular development and impact of a marine eukaryotic parasite on its zombified microalgal host. ISME Journal, 2022, 16, 2348-2359.	4.4	10
64	Redescription and phylogenetic analyses of Durchoniella spp. (Ciliophora, Astomatida) associated with the polychaete Cirriformia tentaculata (Montagu, 1808). European Journal of Protistology, 2017, 61, 265-277.	0.5	9
65	Dinophyceae can use exudates as weapons against the parasite <i>Amoebophrya</i> sp. (Syndiniales). ISME Communications, 2021, 1, .	1.7	8
66	Marine gregarine genomes reveal the breadth of apicomplexan diversity with a partially conserved glideosome machinery. BMC Genomics, 2022, 23, .	1.2	7
67	First report of vampyrellid predator–prey dynamics in a marine system. ISME Journal, 2019, 13, 1110-1113.	4.4	6
68	Dinoflagellate Host Chloroplasts and Mitochondria Remain Functional During Amoebophrya Infection. Frontiers in Microbiology, 2020, 11, 600823.	1.5	6
69	Unveiling protist diversity associated with the Pacific oyster Crassostrea gigas using blocking and excluding primers. BMC Microbiology, 2020, 20, 193.	1.3	6
70	A review of the characteristics of the dinoflagellate parasite Ichthyodinium chabelardi and its potential effect on fin fish populations. Marine and Freshwater Research, 2019, 70, 1307.	0.7	4
71	Temperature Affects the Biological Control of Dinoflagellates by the Generalist Parasitoid Parvilucifera rostrata. Microorganisms, 2022, 10, 385.	1.6	4
72	On the trail of ancestors of diatoms. Journal of Phycology, 2014, 50, 975-976.	1.0	0