

CÃ©dric Berney

List of Publications by Year in descending order

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46
papers

7,541
citations

126907

33
h-index

254184

43
g-index

52
all docs

52
docs citations

52
times ranked

6848
citing authors

#	ARTICLE	IF	CITATIONS
1	Patterns of eukaryotic diversity from the surface to the deep-ocean sediment. <i>Science Advances</i> , 2022, 8, eabj9309.	10.3	52
2	<i>Cyphoderia ampulla</i> (Cyphoderiidae: Rhizaria), a tale of freshwater sailors: The causes and consequences of ecological transitions through the salinity barrier in a family of benthic protists. <i>Molecular Ecology</i> , 2022, 31, 2644-2663.	3.9	7
3	Revisions to the Classification, Nomenclature, and Diversity of Eukaryotes. <i>Journal of Eukaryotic Microbiology</i> , 2019, 66, 4-119.	1.7	904
4	Rhizarian "Novel Clade 10" Revealed as Abundant and Diverse Planktonic and Terrestrial Flagellates, including <i>Aquavolon</i> n. gen.. <i>Journal of Eukaryotic Microbiology</i> , 2018, 65, 828-842.	1.7	29
5	Clarifying the Relationships between Microsporidia and Cryptomycota. <i>Journal of Eukaryotic Microbiology</i> , 2018, 65, 773-782.	1.7	98
6	Environmental Sequencing Fills the Gap Between Parasitic Haplosporidians and Free-living Giant Amoebae. <i>Journal of Eukaryotic Microbiology</i> , 2018, 65, 574-586.	1.7	21
7	EukRef: Phylogenetic curation of ribosomal RNA to enhance understanding of eukaryotic diversity and distribution. <i>PLoS Biology</i> , 2018, 16, e2005849.	5.6	101
8	Soil protistology rebooted: 30 fundamental questions to start with. <i>Soil Biology and Biochemistry</i> , 2017, 111, 94-103.	8.8	130
9	High-throughput sequencing of microbial eukaryotes in Lake Baikal reveals ecologically differentiated communities and novel evolutionary radiations. <i>FEMS Microbiology Ecology</i> , 2017, 93, .	2.7	35
10	<i>UniEuk</i> : Time to Speak a Common Language in Protistology!. <i>Journal of Eukaryotic Microbiology</i> , 2017, 64, 407-411.	1.7	74
11	Parasites dominate hyperdiverse soil protist communities in Neotropical rainforests. <i>Nature Ecology and Evolution</i> , 2017, 1, 91.	7.8	262
12	Phylogeny and Systematics of Leptomyxid Amoebae (Amoebozoa, Tubulinea, Leptomyxida). <i>Protist</i> , 2017, 168, 220-252.	1.5	11
13	A new phylogeny and environmental DNA insight into paramyxids: an increasingly important but enigmatic clade of protistan parasites of marine invertebrates. <i>International Journal for Parasitology</i> , 2016, 46, 605-619.	3.1	39
14	The Large Subunit rDNA Sequence of <i>Plasmodiophora brassicae</i> Does not Contain Intra-species Polymorphism. <i>Protist</i> , 2016, 167, 544-554.	1.5	30
15	Eukaryotic plankton diversity in the sunlit ocean. <i>Science</i> , 2015, 348, 1261605.	12.6	1,551
16	Expansion of the "Reticulosphere": Diversity of Novel Branching and Network-forming Amoebae Helps to Define Variosea (Amoebozoa). <i>Protist</i> , 2015, 166, 271-295.	1.5	57
17	Multigene phylogeny resolves deep branching of Amoebozoa. <i>Molecular Phylogenetics and Evolution</i> , 2015, 83, 293-304.	2.7	84
18	Lineage-specific molecular probing reveals novel diversity and ecological partitioning of haplosporidians. <i>ISME Journal</i> , 2014, 8, 177-186.	9.8	61

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19	Diverse molecular signatures for ribosomally â€œactiveâ€™ Perkinsea in marine sediments. BMC Microbiology, 2014, 14, 110.	3.3	54
20	Mikrocytids Are a Broadly Distributed and Divergent Radiation of Parasites in Aquatic Invertebrates. Current Biology, 2014, 24, 807-812.	3.9	58
21	Multigene eukaryote phylogeny reveals the likely protozoan ancestors of opisthokonts (animals,) Tj ETQq1 1 0.784314 rgBT /Overloc 2.7 97		
22	Vampires in the oceans: predatory cercozoan amoebae in marine habitats. ISME Journal, 2013, 7, 2387-2399.	9.8	73
23	CBOL Protist Working Group: Barcoding Eukaryotic Richness beyond the Animal, Plant, and Fungal Kingdoms. PLoS Biology, 2012, 10, e1001419.	5.6	488
24	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.	14.5	1,463
25	Reticulamoeba Is a Long-Branched Granofilosean (Cercozoa) That Is Missing from Sequence Databases. PLoS ONE, 2012, 7, e49090.	2.5	24
26	The Novel Marine Gliding Zooflagellate Genus Mantamonas (Mantamonadida ord. n.: Apusozoa). Protist, 2011, 162, 207-221.	1.5	49
27	Freshwater Perkinsea and marine-freshwater colonizations revealed by pyrosequencing and phylogeny of environmental rDNA. ISME Journal, 2010, 4, 1144-1153.	9.8	208
28	Phylogeny of Novel Naked Filose and Reticulose Cercozoa: Granofilosea cl. n. and Proteomyxidea Revised. Protist, 2009, 160, 75-109.	1.5	146
29	Diversification of unicellular eukaryotes: cryptomonad colonizations of marine and fresh waters inferred from revised 18S rRNA phylogeny. Environmental Microbiology, 2008, 10, 2635-2644.	3.8	79
30	Molecular comparison of cultivable protozoa from a pristine and a polycyclic aromatic hydrocarbon polluted site. Soil Biology and Biochemistry, 2007, 39, 139-148.	8.8	49
31	Phylogenetic position of Multicilia marina and the evolution of Amoebozoa. International Journal of Systematic and Evolutionary Microbiology, 2006, 56, 1449-1458.	1.7	60
32	A molecular time-scale for eukaryote evolution recalibrated with the continuous microfossil record. Proceedings of the Royal Society B: Biological Sciences, 2006, 273, 1867-1872.	2.6	175
33	Higherâ€™Order Phylogeny of Plasmodial Slime Molds (Myxogastria) Based on Elongation Factor 1â€™A and Small Subunit rRNA Gene Sequences. Journal of Eukaryotic Microbiology, 2005, 52, 201-210.	1.7	84
34	How many novel eukaryotic "kingdoms"? Pitfalls and limitations of environmental DNA surveys. Journal of Eukaryotic Microbiology, 2005, 52, 7S-27S.	1.7	2
35	A molecular perspective on the phylogeny of amoeboid protists. Journal of Eukaryotic Microbiology, 2005, 52, 7S-27S.	1.7	0
36	Phylogeny of lobose amoebae based on actin and small-subunit ribosomal RNA genes. Journal of Eukaryotic Microbiology, 2005, 52, 7S-27S.	1.7	0

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37	The twilight of sun-animalcules. <i>Journal of Eukaryotic Microbiology</i> , 2005, 52, 7S-27S.	1.7	0
38	Small subunit ribosomal RNA sequences of Phaeodarea challenge the monophyly of Haeckel's Radiolaria. <i>Journal of Eukaryotic Microbiology</i> , 2005, 52, 7S-27S.	1.7	1
39	The Testate Lobose Amoebae (Order Arcellinida Kent, 1880) Finally Find their Home within Amoebozoa. <i>Protist</i> , 2005, 156, 191-202.	1.5	78
40	Molecular Phylogeny and Classification of the Lobose Amoebae. <i>Protist</i> , 2005, 156, 129-142.	1.5	99
41	Small-Subunit Ribosomal RNA Gene Sequences of Phaeodarea Challenge the Monophyly of Haeckel's Radiolaria. <i>Protist</i> , 2004, 155, 53-63.	1.5	63
42	How many novel eukaryotic 'kingdoms'? Pitfalls and limitations of environmental DNA surveys. <i>BMC Biology</i> , 2004, 2, 13.	3.8	177
43	The twilight of Heliozoa and rise of Rhizaria, an emerging supergroup of amoeboid eukaryotes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004, 101, 8066-8071.	7.1	227
44	Revised Small Subunit rRNA Analysis Provides Further Evidence that Foraminifera Are Related to Cercozoa. <i>Journal of Molecular Evolution</i> , 2003, 57, S120-S127.	1.8	39
45	Phylogeny of Lobose Amoebae Based on Actin and Small-Subunit Ribosomal RNA Genes. <i>Molecular Biology and Evolution</i> , 2003, 20, 1881-1886.	8.9	89
46	Phylogenetic Position of Dujardin inferred from Nuclear-Encoded Small Subunit Ribosomal DNA. <i>Protist</i> , 2002, 153, 251-260.	1.5	39