The Revised Classification of Eukaryotes

Journal of Eukaryotic Microbiology 59, 429-514 DOI: 10.1111/j.1550-7408.2012.00644.x

Citation Report

#	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	Protein import into the photosynthetic organelles of Paulinella chromatophora and its implications for primary plastid endosymbiosis. Symbiosis, 2012, 58, 99-107.	1.2	15
3	Field et al. Redux EvoDevo, 2013, 4, 5.	1.3	6
4	The Microtubular Cytoskeleton of the Apusomonad Thecamonas, a Sister Lineage to the Opisthokonts. Protist, 2013, 164, 598-621.	0.6	24
5	Inorganic phosphate uptake in Trypanosoma cruzi is coupled to K+ cycling and to active Na+ extrusion. Biochimica Et Biophysica Acta - General Subjects, 2013, 1830, 4265-4273.	1.1	19
6	The new micro-kingdoms of eukaryotes. BMC Biology, 2013, 11, 40.	1.7	27
7	Flagellar apparatus structure of choanocyte in Sycon sp. and its significance for phylogeny of Porifera. Zoomorphology, 2013, 132, 351-357.	0.4	11
8	Lateral Gene Transfer and the Evolution of Photosynthesis in Eukaryotes. , 2013, , 15-53.		0
9	The Non-dictyostelid Sorocarpic Amoebae. , 2013, , 219-242.		16
10	The Mystery of Clade X: Orciraptor gen. nov. and Viridiraptor gen. nov. are Highly Specialised, Algivorous Amoeboflagellates (Glissomonadida, Cercozoa). Protist, 2013, 164, 706-747.	0.6	52
11	Cryptic organelle homology in apicomplexan parasites: insights from evolutionary cell biology. Current Opinion in Microbiology, 2013, 16, 424-431.	2.3	20
12	Two New Marine Ciliates, <i>Caryotricha rariseta</i> n. sp. and <i>Discocephalus pararotatorius</i> n. sp. (Ciliophora, Spirotrichea), with Phylogenetic Analyses Inferred from the Small Subunit <scp>rRNA</scp> Gene Sequences. Journal of Eukaryotic Microbiology, 2013, 60, 388-398.	0.8	13
13	Phagotrophic Protist Diversity in the Groundwater of a Karstified Aquifer – Morphological and Molecular Analysis. Journal of Eukaryotic Microbiology, 2013, 60, 467-479.	0.8	18
14	Targeting apicoplasts in malaria parasites. Expert Opinion on Therapeutic Targets, 2013, 17, 167-177.	1.5	46
15	NIF-type iron-sulfur cluster assembly system is duplicated and distributed in the mitochondria and cytosol of <i>Mastigamoeba balamuthi</i> . Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 7371-7376.	3.3	60
16	A Short Guide to Common Heterotrophic Flagellates of Freshwater Habitats Based on the Morphology of Living Organisms. Protist, 2013, 164, 842-860.	0.6	40
17	Evolution of Dictyostelid Social Amoebas Inferred from the Use of Molecular Tools. , 2013, , 167-182.		2
18	Phylogenomics demonstrates that breviate flagellates are related to opisthokonts and apusomonads. Proceedings of the Roval Society B: Biological Sciences, 2013, 280, 20131755.	1.2	119

#	Article	IF	CITATIONS
19	Giant protists (xenophyophores and komokiaceans) from the Clarion-Clipperton ferromanganese nodule field (eastern Pacific). Biology Bulletin Reviews, 2013, 3, 388-398.	0.3	27
20	The amoeboid protists of cryogenic soils in the Kolyma Lowland. Eurasian Soil Science, 2013, 46, 1211-1218.	0.5	3
21	Microbial Eukaryote Globins. Advances in Microbial Physiology, 2013, 63, 391-446.	1.0	36
22	Scaling body size fluctuations. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 4646-4650.	3.3	77
23	Detection of the kinetoplastid Azumiobodo hoyamushi, the causative agent of soft tunic syndrome, in wild ascidians Halocynthia roretzi. Diseases of Aquatic Organisms, 2013, 106, 267-271.	0.5	12
24	Multicellularity arose several times in the evolution of eukaryotes (Response to DOI) Tj ETQq1 1 0.784314 rgBT	/Overlock 1.2	10 ₅ f 50 54 <mark>2</mark>
25	Evolution of Archamoebae: Morphological and Molecular Evidence for Pelobionts Including Rhizomastix, Entamoeba, Iodamoeba, and Endolimax. Protist, 2013, 164, 380-410.	0.6	42
26	The other eukaryotes in light of evolutionary protistology. Biology and Philosophy, 2013, 28, 299-330.	0.7	20
27	A second rhodopsin-like protein in Cyanophora paradoxa: Gene sequence and protein expression in a cell-free system. Journal of Photochemistry and Photobiology B: Biology, 2013, 125, 188-193.	1.7	3
28	Culture of the cladoceran Moina macrocopa: Mortality associated with flagellate infection. Aquaculture, 2013, 416-417, 374-379.	1.7	8
29	Euglyphid Testate Amoebae (Rhizaria: Euglyphida) from an Arctic Eocene Waterbody: Evidence of Evolutionary Stasis in Plate Morphology For Over 40 Million Years. Protist, 2013, 164, 541-555.	0.6	16
30	Evolution of microtubule organizing centers across the tree of eukaryotes. Plant Journal, 2013, 75, 230-244.	2.8	98
31	Diversity and phylogeny of insect trypanosomatids: all that is hidden shall be revealed. Trends in Parasitology, 2013, 29, 43-52.	1.5	173
32	A New Freshwater Amoeba: <i>Cochliopodium pentatrifurcatum</i> n. sp. (Amoebozoa, Amorphea). Journal of Eukaryotic Microbiology, 2013, 60, 342-349.	0.8	24
33	Chloroplast genome of one brown seaweed, Saccharina japonica (Laminariales, Phaeophyta): Its structural features and phylogenetic analyses with other photosynthetic plastids. Marine Genomics, 2013, 10, 1-9.	0.4	28
34	Algal taxonomy forum. Journal of Phycology, 2013, 49, 226-228.	1.0	2
35	A Modern Descendant of Early Green Algal Phagotrophs. Current Biology, 2013, 23, 1081-1084.	1.8	77
36	Microsporidia and â€~The Art of Living Together'. Advances in Parasitology, 2013, 82, 253-319.	1.4	210

\mathbf{c}		Der	PORT
			ו או אי
<u> </u>	/	IVEI	

#	Article	IF	CITATIONS
37	The Importance of the 45 S Ribosomal Small Subunit-related Complex for Mitochondrial Translation in Trypanosoma brucei. Journal of Biological Chemistry, 2013, 288, 32963-32978.	1.6	24
38	The Ancient and Widespread Nature of the ER–Mitochondria Encounter Structure. Molecular Biology and Evolution, 2013, 30, 2044-2049.	3.5	90
39	Two new genera and species of the monothalamous foraminifera from coastal waters of the Black Sea. Marine Biodiversity, 2013, 43, 473-479.	0.3	7
40	Cell Biology of Chromerids. International Review of Cell and Molecular Biology, 2013, 306, 333-369.	1.6	26
41	Characterization of Amoeboaphelidium protococcarum, an Algal Parasite New to the Cryptomycota Isolated from an Outdoor Algal Pond Used for the Production of Biofuel. PLoS ONE, 2013, 8, e56232.	1.1	136
42	Upper Arctic Ocean water masses harbor distinct communities of heterotrophic flagellates. Biogeosciences, 2013, 10, 4273-4286.	1.3	33
43	Evolution of Tre-2/Bub2/Cdc16 (TBC) Rab GTPase-activating proteins. Molecular Biology of the Cell, 2013, 24, 1574-1583.	0.9	57
44	A genome-wide analysis of annexins from parasitic organisms and their vectors. Scientific Reports, 2013, 3, 2893.	1.6	31
45	Diversity and Distribution of Marine Microbial Eukaryotes. , 2013, , 1-5.		2
46	The CCAP KnowledgeBase: linking protistan and cyanobacterial biological resources with taxonomic and molecular data. Systematics and Biodiversity, 2013, 11, 407-413.	0.5	20
47	Accommodating the load. Mobile Genetic Elements, 2013, 3, e24775.	1.8	30
48	An Advanced System of the Mitochondrial Processing Peptidase and Core Protein Family in Trypanosoma brucei and Multiple Origins of the Core I Subunit in Eukaryotes. Genome Biology and Evolution, 2013, 5, 860-875.	1.1	16
49	Insights into the Origin of Metazoan Filopodia and Microvilli. Molecular Biology and Evolution, 2013, 30, 2013-2023.	3.5	78
50	Evolutionary cell biology of chromosome segregation: insights from trypanosomes. Open Biology, 2013, 3, 130023.	1.5	70
51	A case of taxonomic inflation in coccoid algae: <i>Ellipsoidion parvum</i> and <i>Neocystis vischeri</i> are conspecific with Neocystis (= <i>Nephrodiella</i>) brevis (Chlorophyta,) Tj ETQq0 0 0 rgBT /Overl	oc lo.110 Tf	509177 Td (T
52	Variation of Storage Polysaccharides in Phototrophic Microorganisms. Journal of Applied Glycoscience (1999), 2013, 60, 21-27.	0.3	41
53	Occurrence of homospermidine and thermospermine as a cellular polyamine in unicellular chlorophyte and multicellular charophyte green algae. Journal of General and Applied Microbiology, 2013, 59, 313-319.	0.4	11
54	Picomonas judraskeda Gen. Et Sp. Nov.: The First Identified Member of the Picozoa Phylum Nov., a Widespread Group of Picoeukaryotes, Formerly Known as â€~Picobiliphytes'. PLoS ONE, 2013, 8, e59565.	1.1	97

ARTICLE IF CITATIONS # â€~Candidatus Megaira polyxenophila' gen. nov., sp. nov.: Considerations on Evolutionary History, Host 55 1.1 76 Range and Shift of Early Divergent Rickettsiae. PLoS ONE, 2013, 8, e72581. Evolution and Distribution of Saxitoxin Biosynthesis in Dinoflagellates. Marine Drugs, 2013, 11, 2.2 58 2814-2828. Phytoplankton of the northern coastal and shelf waters of the Yucatan Peninsula, southeastern Gulf 57 0.1 19 of Mexico, Mexico. Check List, 2013, 9, 771. Description of Colponema vietnamica sp.n. and Acavomonas peruviana n. gen. n. sp., Two New Alveolate Phyla (Colponemidia nom. nov. and Acavomonidia nom. nov.) and Their Contributions to Reconstructing the Ancestral State of Alveolates and Eukaryotes. PLoS ONE, 2014, 9, e95467. Phylogenomic Analysis of "Red―Genes from Two Divergent Species of the "Green―Secondary Phototrophs, the Chlorarachniophytes, Suggests Multiple Horizontal Gene Transfers from the Red 59 1.1 9 Lineage before the Divergence of Extant Chlorarachniophytes. PLoS ONE, 2014, 9, e101158. Primary endosymbiosis and the evolution of light and oxygen sensing in photosynthetic eukaryotes. 1.1 Frontiers in Ecology and Evolution, 2014, 2, . Local factors affecting the testate amoeba community (Protozoa: Arcellinida; Euglyphida) in a 62 0.3 8 neotropical floodplain. Journal of Limnology, 2014, 73, . Evolution and Classification of Myosins, a Paneukaryotic Whole-Genome Approach. Genome Biology 1.1 121 and Evolution, 2014, 6, 290-305. 64 A Trip Through Chemical Space: Why Life Has Evolved the Chemistry That It Has., 2014, , 371-394. 1 Fragmentation of tRNA in Phytophthora infestans asexual life cycle stages and during host plant 1.3 24 infection. BMC Microbiology, 2014, 14, 308. Vaccines against bovine babesiosis: where we are now and possible roads ahead. Parasitology, 2014, 141, 0.7 70 66 1563-1592. Exploring the evolutionary history of centrosomes. Philosophical Transactions of the Royal Society 1.8 B: Biological Sciences, 2014, 369, 20130453. Communities of microbial eukaryotes in the mammalian gut within the context of environmental 68 1.5 130 eukaryotic diversity. Frontiers in Microbiology, 2014, 5, 298. Ecological functions of zoosporic hyperparasites. Frontiers in Microbiology, 2014, 5, 244. 69 1.5 Dynamics and drivers of the protozoic Si pool along a 10-year chronosequence of initial ecosystem 70 1.6 38 states. Ecological Engineering, 2014, 70, 477-482. Widespread occurrence of organelle genome-encoded 5S rRNAs including permuted molecules. Nucleic Acids Research, 2014, 42, 13764-13777. 129 The Genome of Spironucleus salmonicida Highlights a Fish Pathogen Adapted to Fluctuating 72 1.563 Environments. PLoS Genetics, 2014, 10, e1004053. Microbiotic signatures of the Anthropocene in marginal marine and freshwater palaeoenvironments. Geological Society Special Publication, 2014, 395, 185-219.

#	ARTICLE Bayesian modelling of compositional heterogeneity in molecular phylogenetics. Statistical	IF 0.2	CITATIONS
74 76	Applications in Genetics and Molecular Biology, 2014, 13, 589-609. The evolution of photosynthesis in chromist algae through serial endosymbioses. Nature Communications, 2014, 5, 5764.	5.8	130
78	Five <i><scp>C</scp>yanophora</i> (<scp>C</scp> yanophorales, <scp>G</scp> laucophyta) species delineated based on morphological and molecular data. Journal of Phycology, 2014, 50, 1058-1069.	1.0	18
79	Missing Genes, Multiple ORFs, and C-to-U Type RNA Editing in Acrasis kona (Heterolobosea, Excavata) Mitochondrial DNA. Genome Biology and Evolution, 2014, 6, 2240-2257.	1.1	26
80	Distribution of free-living amoebae in a treatment system of textile industrial wastewater. Experimental Parasitology, 2014, 145, S34-S38.	0.5	25
81	Soil water availability strongly alters the community composition of soil protists. Pedobiologia, 2014, 57, 205-213.	0.5	125
82	Global diversity and geography of soil fungi. Science, 2014, 346, 1256688.	6.0	2,513
83	Phylogeny of the Poorly Known Ciliates, Microthoracida, a Systematically Confused Taxon (Ciliophora), with Morphological Reports of Three Species. Journal of Eukaryotic Microbiology, 2014, 61, 227-237.	0.8	10
84	Ribosomal Gene Polymorphism in Small Genomes: Analysis of Different 16S <scp>rRNA</scp> Sequences Expressed in the Honeybee Parasite <i>Nosema ceranae</i> (Microsporidia). Journal of Eukaryotic Microbiology, 2014, 61, 42-50.	0.8	11
85	Interactions of Foodborne Pathogens with Freeâ€ŀiving Protozoa: Potential Consequences for Food Safety. Comprehensive Reviews in Food Science and Food Safety, 2014, 13, 924-944.	5.9	34
86	Chromera velia, Endosymbioses and the Rhodoplex Hypothesis—Plastid Evolution in Cryptophytes, Alveolates, Stramenopiles, and Haptophytes (CASH Lineages). Genome Biology and Evolution, 2014, 6, 666-684.	1.1	93
87	Evolution of the Cytosolic Iron-Sulfur Cluster Assembly Machinery in Blastocystis Species and Other Microbial Eukaryotes. Eukaryotic Cell, 2014, 13, 143-153.	3.4	47
89	Distribution of Conventional and Nonconventional Introns in Tubulin (α and β) Genes of Euglenids. Molecular Biology and Evolution, 2014, 31, 584-593.	3.5	20
90	Monophyly of Archaeplastida supergroup and relationships among its lineages in the light of phylogenetic and phylogenomic studies. Are we close to a consensus?. Acta Societatis Botanicorum Poloniae, 2014, 83, 263-280.	0.8	27
91	A contemplation on the secondary origin of green algal and plant plastids. Acta Societatis Botanicorum Poloniae, 2014, 83, 331-336.	0.8	13
92	Primary endosymbiosis: have cyanobacteria and Chlamydiae ever been roommates?. Acta Societatis Botanicorum Poloniae, 2014, 83, 291-302.	0.8	23
93	Contrasting patterns in the evolution of the Rab GTPase family in Archaeplastida. Acta Societatis Botanicorum Poloniae, 2014, 83, 303-315.	0.8	15
94	A phylogenetic reconsideration of suctorian ciliates (<scp>P</scp> rotista, <scp>C</scp> iliophora,) Tj ETQq1 1 C Scripta, 2014, 43, 206-216.).784314 ı 0.7	rgBT /Overloci 8

#	Article	IF	CITATIONS
95	Large-scale phylogenomic analysis reveals the phylogenetic position of the problematic taxon Protocruzia and unravels the deep phylogenetic affinities of the ciliate lineages. Molecular Phylogenetics and Evolution, 2014, 78, 36-42.	1.2	71
96	Evolutionary mechanisms for establishing eukaryotic cellular complexity. Trends in Cell Biology, 2014, 24, 435-442.	3.6	26
97	Molecular diversity of endosymbiotic Nephroselmis (Nephroselmidophyceae) in Hatena arenicola (Katablepharidophycota). Journal of Plant Research, 2014, 127, 241-247.	1.2	12
98	Paleobiological Perspectives on Early Eukaryotic Evolution. Cold Spring Harbor Perspectives in Biology, 2014, 6, a016121-a016121.	2.3	298
99	Evolution: Rooting the Eukaryotic Tree of Life. Current Biology, 2014, 24, R151-R152.	1.8	13
100	Nuclear DNA replication initiation in kinetoplastid parasites: new insights into an ancient process. Trends in Parasitology, 2014, 30, 27-36.	1.5	32
101	An Alternative Root for the Eukaryote Tree of Life. Current Biology, 2014, 24, 465-470.	1.8	196
102	Evolution of the nucleus. Current Opinion in Cell Biology, 2014, 28, 8-15.	2.6	49
103	Amplification primers of SSU rDNA for soil protists. Soil Biology and Biochemistry, 2014, 69, 328-342.	4.2	54
104	Pheromone signaling during sexual reproduction in algae. Plant Journal, 2014, 79, 632-644.	2.8	72
105	Kinase signalling in Plasmodium sexual stages and interventions to stop malaria transmission. Molecular and Biochemical Parasitology, 2014, 193, 23-32.	0.5	11
106	The others: our biased perspective of eukaryotic genomes. Trends in Ecology and Evolution, 2014, 29, 252-259.	4.2	167
108	Les amibes libres: un danger méconnu. Revue Francophone Des Laboratoires, 2014, 2014, 41-51.	0.0	3
109	Molecular markers from different genomic compartments reveal cryptic diversity within glaucophyte species. Molecular Phylogenetics and Evolution, 2014, 76, 181-188.	1.2	21
110	A comparative analysis of trypanosomatid SNARE proteins. Parasitology International, 2014, 63, 341-348.	0.6	17
111	The Genome of the Foraminiferan Reticulomyxa filosa. Current Biology, 2014, 24, 11-18.	1.8	73
112	Further insights into the phylogeny of two ciliate classes Nassophorea and Prostomatea (Protista,) Tj ETQq0 0 0	rgBT /Over 1.2	lock 10 Tf 50

113	The diversity and phylogeny of the commercially important algal class Eustigmatophyceae, including the new clade Goniochloridales. Journal of Applied Phycology, 2014, 26, 1773-1782.	1.5	41
-----	---	-----	----

		CITATION REPORT		
#	Article		IF	CITATIONS
114	Ancient signals: comparative genomics of green plant CDPKs. Trends in Plant Science,	2014, 19, 79-89.	4.3	152
115	Paravannella minima n. g. n. sp. (Discosea, Vannellidae) and distinction of the genera ir amoebae. European Journal of Protistology, 2014, 50, 258-269.	n the vannellid	0.5	16
116	The Weng'an biota and the Ediacaran radiation of multicellular eukaryotes. National Sc 2014, 1, 498-520.	cience Review,	4.6	117
117	The SILVA and "All-species Living Tree Project (LTP)―taxonomic frameworks. Nucle 2014, 42, D643-D648.	eic Acids Research,	6.5	2,667
118	Cryptic Sex in <i>Symbiodinium</i> (Alveolata, Dinoflagellata) is Supported by an Inve Genes. Journal of Eukaryotic Microbiology, 2014, 61, 322-327.	ntory of Meiotic	0.8	72
119	Phylogeny, Ultrastructure, and Flagellar Apparatus of a New Marimonad Flagellate Abo sp. nov. (Imbricatea, Cercozoa). Protist, 2014, 165, 808-824.	llifer globosa	0.6	9
120	Complex Patterns of Gene Fission in the Eukaryotic Folate Biosynthesis Pathway. Geno Evolution, 2014, 6, 2709-2720.	me Biology and	1.1	12
122	The Diatom Attachment Scar <i>Ophthalmichnus lyolithon</i> igen. et isp. n Ichnos, 20	014, 21, 111-118.	0.8	10
123	Placing Environmental Next-Generation Sequencing Amplicons from Microbial Eukaryo Phylogenetic Context. Molecular Biology and Evolution, 2014, 31, 993-1009.	tes into a	3.5	97
124	3 Systematics of the Straminipila: Labyrinthulomycota, Hyphochytriomycota, and Oom 39-97.	ıycota. , 2014, ,		56
125	How discordant morphological and molecular evolution among microorganisms can re notions of biodiversity on Earth. BioEssays, 2014, 36, 950-959.	vise our	1.2	36
126	Green Algae and the Origins of Multicellularity in the Plant Kingdom. Cold Spring Harbo Perspectives in Biology, 2014, 6, a016170-a016170.	br	2.3	111
127	Response of the protozooplankton assemblage during the European Iron Fertilization I (EIFEX) in the Antarctic circumpolar current. Journal of Plankton Research, 2014, 36, 1		0.8	9
128	Missing Pieces of an Ancient Puzzle: Evolution of the Eukaryotic Membrane-Trafficking Spring Harbor Perspectives in Biology, 2014, 6, a016048-a016048.	System. Cold	2.3	60
130	Horizontal Gene Transfer in Eukaryotic Plant Pathogens. Annual Review of Phytopathol 583-614.	ogy, 2014, 52,	3.5	126
131	What is the importance of zoonotic trichomonads for human health?. Trends in Parasit 333-341.	tology, 2014, 30,	1.5	92
132	Nucleotide substitution analyses of the glaucophyte Cyanophora suggest an ancestral mutation rate in plastid vs mitochondrial DNA for the Archaeplastida. Molecular Phylog Evolution, 2014, 79, 380-384.	ly lower genetics and	1.2	14
133	Trends in research of antitrypanosomal agents among synthetic heterocycles. Europea Medicinal Chemistry, 2014, 85, 51-64.	n Journal of	2.6	40

#	Article	IF	CITATIONS
134	Gregarine site-heterogeneous 18S rDNA trees, revision of gregarine higher classification, and the evolutionary diversification of Sporozoa. European Journal of Protistology, 2014, 50, 472-495.	0.5	103
135	The effect of inter-annual Atlantic water inflow variability on the planktonic protist community structure in the West Spitsbergen waters during the summer. Journal of Plankton Research, 2014, 36, 1190-1203.	0.8	41
136	The reduced kinome of Ostreococcus tauri: core eukaryotic signalling components in a tractable model species. BMC Genomics, 2014, 15, 640.	1.2	18
137	Six Subgroups and Extensive Recent Duplications Characterize the Evolution of the Eukaryotic Tubulin Protein Family. Genome Biology and Evolution, 2014, 6, 2274-2288.	1.1	110
138	Applications of next-generation sequencing to unravelling the evolutionary history of algae. International Journal of Systematic and Evolutionary Microbiology, 2014, 64, 333-345.	0.8	48
139	Exploring the uncultured microeukaryote majority in the oceans: reevaluation of ribogroups within stramenopiles. ISME Journal, 2014, 8, 854-866.	4.4	157
140	The Cell Biology of the Endocytic System from an Evolutionary Perspective. Cold Spring Harbor Perspectives in Biology, 2014, 6, a016998-a016998.	2.3	34
141	The Eukaryotic Tree of Life from a Global Phylogenomic Perspective. Cold Spring Harbor Perspectives in Biology, 2014, 6, a016147-a016147.	2.3	272
142	Molecular diversity reveals previously undetected air-dispersed protist colonists in a Mediterranean area. Science of the Total Environment, 2014, 478, 70-79.	3.9	21
143	Detailed Process of Shell Construction in the Photosynthetic Testate Amoeba <i>Paulinella chromatophora</i> (Euglyphid, Rhizaria). Journal of Eukaryotic Microbiology, 2014, 61, 317-321.	0.8	16
144	A multilocus timescale for oomycete evolution estimated under three distinct molecular clock models. BMC Evolutionary Biology, 2014, 14, 101.	3.2	53
145	Analysis of EST data of the marine protist Oxyrrhis marina, an emerging model for alveolate biology and evolution. BMC Genomics, 2014, 15, 122.	1.2	26
146	eIF4F-like complexes formed by cap-binding homolog TbEIF4E5 with TbEIF4G1 or TbEIF4G2 are implicated in post-transcriptional regulation in <i>Trypanosoma brucei</i> . Rna, 2014, 20, 1272-1286.	1.6	48
147	Mitochondrial Genome Sequences and Comparative Genomics of <i>Achlya hypogyna</i> and <i>Thraustotheca clavata</i> . Journal of Eukaryotic Microbiology, 2014, 61, 146-154.	0.8	12
148	Co-occurrence of free-living protozoa and foodborne pathogens on dishcloths: Implications for food safety. International Journal of Food Microbiology, 2014, 191, 89-96.	2.1	24
149	Multigene eukaryote phylogeny reveals the likely protozoan ancestors of opisthokonts (animals,) Tj ETQq1 1 0	.784314 rgl 1.2	BT Qverlock
150	<i>Ptolemeba</i> n. gen., a Novel Genus of Hartmannellid Amoebae (Tubulinea, Amoebozoa); with an Emphasis on the Taxonomy of <i>Saccamoeba</i> . Journal of Eukaryotic Microbiology, 2014, 61, 611-619.	0.8	5
151	Effects of longâ€ŧerm differential fertilization on eukaryotic microbial communities in an arable soil: a multiple barcoding approach. Molecular Ecology, 2014, 23, 3341-3355.	2.0	163

#	Article	IF	CITATIONS
152	Evidence for two different morphotypes of Difflugia tuberspinifera from China. European Journal of Protistology, 2014, 50, 205-211.	0.5	9
153	Ultrastructural morphology of the reproductive swarmers of Sphaerozoum punctatum (Huxley) from the East China Sea. European Journal of Protistology, 2014, 50, 194-204.	0.5	10
154	Haplomyxa saranae gen. nov. et sp. nov., a New Naked Freshwater Foraminifer. Protist, 2014, 165, 317-329.	0.6	11
155	Phylogeny and Ultrastructure of Oxymonas jouteli, a Rostellum-free Species, and Opisthomitus longiflagellatus sp. nov., Oxymonadid Flagellates from the Gut of Neotermes jouteli. Protist, 2014, 165, 384-399.	0.6	11
156	Phylogenomic analyses support the bifurcation of ciliates into two major clades that differ in properties of nuclear division. Molecular Phylogenetics and Evolution, 2014, 70, 240-243.	1.2	33
157	The predicted secretomes of Monosiga brevicollis and Capsaspora owczarzaki, close unicellular relatives of metazoans, reveal new insights into the evolution of the metazoan extracellular matrix. Matrix Biology, 2014, 37, 60-68.	1.5	27
158	The <scp>D</scp> 1â€ <scp>D</scp> 2 region of the large subunit ribosomal <scp>DNA</scp> as barcode for ciliates. Molecular Ecology Resources, 2014, 14, 458-468.	2.2	52
159	Effects of light and prey availability on Arctic freshwater protist communities examined by high-throughput DNA and RNA sequencing. FEMS Microbiology Ecology, 2014, 88, 550-564.	1.3	62
160	The rise and fall of Picobiliphytes: How assumed autotrophs turned out to be heterotrophs. BioEssays, 2014, 36, 468-474.	1.2	31
161	Nuclear pore complex evolution: a trypanosome Mlp analogue functions in chromosomal segregation but lacks transcriptional barrier activity. Molecular Biology of the Cell, 2014, 25, 1421-1436.	0.9	26
162	DNA Barcoding in Amoebozoa and Challenges: The Example of Cochliopodium. Protist, 2014, 165, 473-484.	0.6	29
163	Creneis carolina gen. et sp. nov. (Heterolobosea), a Novel Marine Anaerobic Protist with Strikingly Derived Morphology and Life Cycle. Protist, 2014, 165, 542-567.	0.6	21
164	Evolution of parasitism in kinetoplastid flagellates. Molecular and Biochemical Parasitology, 2014, 195, 115-122.	0.5	200
165	Focusing on Genera to Improve Species Identification: Revised Systematics of the Ciliate Spirostomum. Protist, 2014, 165, 527-541.	0.6	37
166	Comparative Ultrastructure and Molecular Phylogeny of Selenidium melongena n. sp. and S. terebellae Ray 1930 Demonstrate Niche Partitioning in Marine Gregarine Parasites (Apicomplexa). Protist, 2014, 165, 493-511.	0.6	20
167	Osmotolerance in the Cryptophyceae: Jacks-of-all-trades in the Chroomonas Clade. Protist, 2014, 165, 123-143.	0.6	21
168	The mitochondrial respiratory chain of the secondary green alga Euglena gracilis shares many additional subunits with parasitic Trypanosomatidae. Mitochondrion, 2014, 19, 338-349.	1.6	59
169	Ciliate Diversity, Community Structure, and Novel Taxa in Lakes of the McMurdo Dry Valleys, Antarctica. Biological Bulletin, 2014, 227, 175-190.	0.7	15

#	Article	IF	CITATIONS
170	General Overview. , 2014, , 1-48.		1
171	Identification of the meiotic toolkit in diatoms and exploration of meiosis-specific SPO11 and RAD51 homologs in the sexual species Pseudo-nitzschia multistriata and Seminavis robusta. BMC Genomics, 2015, 16, 930.	1.2	53
172	Seasonal dynamics of culturable thraustochytrids (Labyrinthulomycetes, Stramenopiles) in estuarine and coastal waters. Aquatic Microbial Ecology, 2015, 74, 187-204.	0.9	44
173	Polyphyly of nuclear lamin genes indicates an early eukaryotic origin of the metazoan-type intermediate filament proteins. Scientific Reports, 2015, 5, 10652.	1.6	39
174	Marine amoebae with cytoplasmic and perinuclear symbionts deeply branching in the Gammaproteobacteria. Scientific Reports, 2015, 5, 13381.	1.6	36
175	Heterotrophic Planktonic Microbes: Virus, Bacteria, Archaea, and Protozoa. , 2015, , 4.2.2-1-4.2.2-34.		8
176	Testate Amoebae Diversity, from the Atlantic Forest Aquatic and Edaphic Environments, Collected Within the Rio de Janeiro State, Brazil. Microscopy and Microanalysis, 2015, 21, 931-932.	0.2	0
177	Ultra-high voltage electron microscopy of primitive algae illuminates 3D ultrastructures of the first photosynthetic eukaryote. Scientific Reports, 2015, 5, 14735.	1.6	8
178	Analysis of the dinoflagellate Prorocentrum minimum transcriptome: Identifying the members of the voltage-gated cation channel superfamily. Cell and Tissue Biology, 2015, 9, 483-492.	0.2	7
179	First record of siliceous shell plates of Scutiglypha sp. (Amoebozoa: Thecamoeba: Euglyphidae) from Lake Enspel (upper Oligocene, Westerwald Mountains, Germany). Palaeobiodiversity and Palaeoenvironments, 2015, 95, 513-519.	0.6	2
180	Evolution of the Germline Actin Gene in Hypotrichous Ciliates: Multiple Nonscrambled <scp>IES</scp> s at Extremely Conserved Locations in Two Urostylids. Journal of Eukaryotic Microbiology, 2015, 62, 188-195.	0.8	4
181	Evolutionary history of phosphatidylinositol- 3-kinases: ancestral origin in eukaryotes and complex duplication patterns. BMC Evolutionary Biology, 2015, 15, 226.	3.2	16
182	Analysis of the mitochondrial maxicircle of Trypanosoma lewisi, a neglected human pathogen. Parasites and Vectors, 2015, 8, 665.	1.0	27
183	Characterization of Two Species of Trypanosomatidae from the Honey Bee <i>Apis mellifera</i> : <i>Crithidia mellificae</i> Langridge and McGhee, and <i>Lotmaria passim</i> n. gen., n. sp Journal of Eukaryotic Microbiology, 2015, 62, 567-583.	0.8	152
184	Not all are freeâ€living: highâ€throughput <scp>DNA</scp> metabarcoding reveals a diverse community of protists parasitizing soil metazoa. Molecular Ecology, 2015, 24, 4556-4569.	2.0	116
185	Morphology and molecular phylogeny of three new oligotrich ciliates (Protozoa, Ciliophora) from the South China Sea. Zoological Journal of the Linnean Society, 2015, 174, 653-665.	1.0	19
186	Sterol Composition and Biosynthetic Genes of <i>Vitrella brassicaformis</i> , a Recently Discovered Chromerid: Comparison to <i>Chromera velia</i> and Phylogenetic Relationship with Apicomplexan Parasites. Journal of Eukaryotic Microbiology, 2015, 62, 786-798.	0.8	2
187	Marine protist diversity in <scp>E</scp> uropean coastal waters and sediments as revealed by highâ€ŧhroughput sequencing. Environmental Microbiology, 2015, 17, 4035-4049.	1.8	384

#	Article	IF	CITATIONS
188	Cultivation and Characterisation of New Species of Apusomonads (the Sister Group to Opisthokonts), Including Close Relatives of <i>Thecamonas</i> (<i>Chelonemonas</i> n. gen.). Journal of Eukaryotic Microbiology, 2015, 62, 637-649.	0.8	9
189	Pack hunting by a common soil amoeba on nematodes. Environmental Microbiology, 2015, 17, 4538-4546.	1.8	93
190	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
191	<i>Cochliopodium arabianum</i> n. sp. (Amorphea, Amoebozoa). Journal of Eukaryotic Microbiology, 2015, 62, 623-628.	0.8	9
192	The diversity of algal phospholipase D homologs revealed by biocomputational analysis. Journal of Phycology, 2015, 51, 943-962.	1.0	13
193	Effect of daily variation on the seasonality of testate amoebae community structure in different environments of a Neotropical floodplain. Journal of Limnology, 2015, , .	0.3	1
194	Exploring the environmental diversity of kinetoplastid flagellates in the high-throughput DNA sequencing era. Memorias Do Instituto Oswaldo Cruz, 2015, 110, 956-965.	0.8	75
195	Testate amoebae (Protozoa Rhizopoda) in two biotopes of Ubatiba stream, MaricÃį, Rio de Janeiro State. Acta Scientiarum - Biological Sciences, 2015, 37, 291.	0.3	2
196	Form, Fabric, and Function of a Flagellum-Associated Cytoskeletal Structure. Cells, 2015, 4, 726-747.	1.8	20
197	A checklist of ciliate parasites (Ciliophora) of fishes from Mexico. Zootaxa, 2015, 4027, 270-80.	0.2	7
198	Clinical and epidemiological aspects of feline leishmaniasis in Brazil. Semina:Ciencias Agrarias, 2015, 36, 1467.	0.1	9
199	Ancient Origin of the U2 Small Nuclear RNA Gene-Targeting Non-LTR Retrotransposons Utopia. PLoS ONE, 2015, 10, e0140084.	1.1	8
200	A Higher Level Classification of All Living Organisms. PLoS ONE, 2015, 10, e0119248.	1.1	298
201	Ancient Homology of the Mitochondrial Contact Site and Cristae Organizing System Points to an Endosymbiotic Origin of Mitochondrial Cristae. Current Biology, 2015, 25, 1489-1495.	1.8	95
202	The ancient roots of calcium signalling evolutionary tree. Cell Calcium, 2015, 57, 123-132.	1.1	74
203	RÃ1e des animaux vertébrés dans la transmission des champignons dermatophytes pathogènes pour l'homme. Revue Francophone Des Laboratoires, 2015, 2015, 53-60.	0.0	0
204	Identification of the ISWI Chromatin Remodeling Complex of the Early Branching Eukaryote Trypanosoma brucei. Journal of Biological Chemistry, 2015, 290, 26954-26967.	1.6	21
205	Evolutionary cell biology: functional insight from "endless forms most beautiful― Molecular Biology of the Cell, 2015, 26, 4532-4538.	0.9	17

#	Article	IF	CITATIONS
206	Unexpectedly Streamlined Mitochondrial Genome of the Euglenozoan <i>Euglena gracilis</i> . Genome Biology and Evolution, 2015, 7, 3358-3367.	1.1	57
207	Phylogenomics in Algal Research: Current Trends and Future Perspectives. Cellular Origin and Life in Extreme Habitats, 2015, , 501-517.	0.3	1
208	Unusual Features of Dinokaryon, the Enigmatic Nucleus of Dinoflagellates. , 2015, , 23-45.		4
209	Diversity and Ecology of Thraustochytrid Protists in the Marine Environment. , 2015, , 331-346.		11
210	The mineral composition of the tests of â€~testate amoebae' (Amoebozoa, Arcellinida): The relative importance of grain availability and grain selection. Revue De Micropaleontologie, 2015, 58, 141-154.	0.8	11
211	Morphology and life cycle of amoeboflagellate Pharyngomonas sp. (Heterolobosea, Excavata) from hypersaline inland Razval Lake. Biology Bulletin, 2015, 42, 759-769.	0.1	4
212	Morphological and Molecular Evidence Support a Close Relationship Between the Free-living Archamoebae Mastigella and Pelomyxa. Protist, 2015, 166, 14-41.	0.6	23
213	Taxon-Rich Phylogenomic Analyses Resolve the Eukaryotic Tree of Life and Reveal the Power of Subsampling by Sites. Systematic Biology, 2015, 64, 406-415.	2.7	63
214	First report of Enterocytozoon bieneusi in pigs in Brazil. Parasitology International, 2015, 64, 18-23.	0.6	42
215	Rethinking the marine carbon cycle: Factoring in the multifarious lifestyles of microbes. Science, 2015, 347, 1257594.	6.0	679
216	Role of the Cytosolic Heat Shock Protein 70 Ssa5 in the Ciliate Protozoan <i>Tetrahymena thermophila</i> . Journal of Eukaryotic Microbiology, 2015, 62, 481-493.	0.8	7
217	A review of Theileria diagnostics and epidemiology. International Journal for Parasitology: Parasites and Wildlife, 2015, 4, 104-118.	0.6	196
218	Lineageâ€specific activities of a multipotent mitochondrion of trypanosomatid flagellates. Molecular Microbiology, 2015, 96, 55-67.	1.2	12
219	A molecular survey of freshwater microeukaryotes in an Arctic reservoir (Svalbard, 79°N) in summer by using next-generation sequencing. Polar Biology, 2015, 38, 179-187.	0.5	10
220	Bacterial proteins pinpoint a single eukaryotic root. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, E693-9.	3.3	159
221	Conservation of an intricate circuit for crucial modifications of the tRNA ^{Phe} anticodon loop in eukaryotes. Rna, 2015, 21, 61-74.	1.6	60
222	Identification of Parabodo caudatus (class Kinetoplastea) in urine voided from a dog with hematuria. Journal of Veterinary Diagnostic Investigation, 2015, 27, 117-120.	0.5	8
223	Update of information on perkinsosis in NW Mediterranean coast: Identification of Perkinsus spp. (Protista) in new locations and hosts. Journal of Invertebrate Pathology, 2015, 125, 37-41.	1.5	16

ARTICLE

CITATION REPORT IF CITATIONS Free-Living Protozoa., 2015, , 113-132. 6 224 Early evolution of the Eukaryota. Palaeontology, 2015, 58, 5-17. 1.0 The Plastid Terminal Oxidase: Its Elusive Function Points to Multiple Contributions to Plastid 226 8.6 147 Physiology. Annual Review of Plant Biology, 2015, 66, 49-74. EARLY EVOLUTION AND NEW CLASSIFICATION OF THE ORDER ROBERTINIDA (FORAMINIFERA). Journal of 0.1 Foraminiferal Research, 2015, 45, 3-28. The Eukaryotic Ancestor Had a Complex Ubiquitin Signaling System of Archaeal Origin. Molecular 228 3.5 58 Biology and Evolution, 2015, 32, 726-739. Continuous monitoring of near-bottom mesoplankton communities in the East China Sea during a 229 series of typhoons. Journal of Oceanography, 2015, 71, 115-124. Stochastic developmental variation, an epigenetic source of phenotypic diversity with far-reaching 230 0.5 118 biological consequences. Journal of Biosciences, 2015, 40, 159-204. Arcella peruviana sp. nov. (Amoebozoa: Arcellinida, Arcellidae), a new species from a tropical peatland in Amażonia. European Journal of Protistology, 2015, 51, 437-449. 232 Diverse protist grazers select for virulence-related traits in Legionella. ISME Journal, 2015, 9, 1607-1618. 4.4 52 Malleable Mitochondrion of Trypanosoma brucei. International Review of Cell and Molecular 1.6 Biology, 2015, 315, 73-151. Diversity and distribution of unicellular opisthokonts along the <scp>E</scp>uropean coast analysed 234 1.8 52 using highâ€throughput sequencing. Environmental Microbiology, 2015, 17, 3195-3207. Efficient tryptophanâ€catabolizing activity is consistently conserved through evolution of TDO enzymes, but not IDO enzymes. Journal of Experimental Zoology Part B: Molecular and Developmental Evolution, 2015, 324, 128-140. Ancient dynamin segments capture early stages of hostâ€"mitochondrial integration. Proceedings of 236 3.3 41 the National Academy of Sciences of the United States of America, 2015, 112, 2800-2805.

Cu(II) pollution affects fecundity of the mangrove degrader community, the Labyrinthulomycetes. Botanica Marina, 2015, 58, 129-138. 0.6 Perkinsosis in the clams Ruditapes decussatus and R. philippinarum in the Northeastern Atlantic and 238 32 1.5 Mediterranean Sea: A review. Journal of Invertebrate Pathology, 2015, 131, 58-67. ITS2 Database V: Twice as Much: Table 1.. Molecular Biology and Evolution, 2015, 32, 3030-3032. 231

240	The protozoic Si pool in temperate forest ecosystems — Quantification, abiotic controls and interactions with earthworms. Geoderma, 2015, 243-244, 196-204.	2.3	65
241	Liaisons dangereuses: sexual recombination among pathogenic trypanosomes. Research in Microbiology, 2015, 166, 459-466.	1.0	24

# 242	ARTICLE Insights into thermoadaptation and the evolution of mesophily from the bacterial phylum <i>Thermotogae</i> . Canadian Journal of Microbiology, 2015, 61, 655-670.	IF 0.8	CITATIONS
243	Sex is a ubiquitous, ancient, and inherent attribute of eukaryotic life. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 8827-8834.	3.3	236
244	Heterogeneity in the genus Allovahlkampfia and the description of the new genus Parafumarolamoeba (Vahlkampfiidae; Heterolobosea). European Journal of Protistology, 2015, 51, 335-349.	0.5	14
245	Checklist, diversity and distribution of testate amoebae in Chile. European Journal of Protistology, 2015, 51, 409-424.	0.5	44
246	Phylogenetic framework of the systematically confused Anteholosticha–Holosticha complex (Ciliophora, Hypotrichia) based on multigene analysis. Molecular Phylogenetics and Evolution, 2015, 91, 238-247.	1.2	45
247	Gamete Dialogs in Green Lineages. Molecular Plant, 2015, 8, 1442-1454.	3.9	32
248	The Secreted Proteins of Achlya hypogyna and Thraustotheca clavata Identify the Ancestral Oomycete Secretome and Reveal Gene Acquisitions by Horizontal Gene Transfer. Genome Biology and Evolution, 2015, 7, 120-135.	1.1	34
249	The Organellar Genomes of <i>Chromera</i> and <i>Vitrella</i> , the Phototrophic Relatives of Apicomplexan Parasites. Annual Review of Microbiology, 2015, 69, 129-144.	2.9	58
250	Chrysophyceae and Phaeothamniophyceae. , 2015, , 537-586.		18
251	Dinoflagellates. , 2015, , 773-807.		15
252	High diversity of protistan plankton communities in remote high mountain lakes in the European Alps and the Himalayan mountains. FEMS Microbiology Ecology, 2015, 91, .	1.3	61
254	Ciliates $\hat{a} \in$ Protists with complex morphologies and ambiguous early fossil record. Marine Micropaleontology, 2015, 119, 1-6.	0.5	17
255	Expansion of the â€~Reticulosphere': Diversity of Novel Branching and Network-forming Amoebae Helps to Define Variosea (Amoebozoa). Protist, 2015, 166, 271-295.	0.6	57
256	Molecular Phylogeny of the Widely Distributed Marine Protists, Phaeodaria (Rhizaria, Cercozoa). Protist, 2015, 166, 363-373.	0.6	24
257	The past, present and future of mitochondrial genomics: have we sequenced enough mtDNAs?. Briefings in Functional Genomics, 2016, 15, elv027.	1.3	83
258	Infections by Intestinal Coccidia and Giardia duodenalis. Clinics in Laboratory Medicine, 2015, 35, 423-444.	0.7	74
259	Phylogenomics of non-model ciliates based on transcriptomic analyses. Protein and Cell, 2015, 6, 373-385.	4.8	31
260	From Environmental Sequences to Morphology: Observation and Characterisation of a Paulinellid Testate Amoeba (Micropyxidiella edaphonis gen. nov. sp. nov. Euglyphida, Paulinellidae) from Soil using Fluorescent in situ Hybridization. Protist, 2015, 166, 264-270.	0.6	12

#	Article	IF	CITATIONS
261	The two-domain tree of life is linked to a new root for the Archaea. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 6670-6675.	3.3	234
262	Evolution of centrosomes and the nuclear lamina: Amoebozoan assets. European Journal of Cell Biology, 2015, 94, 249-256.	1.6	37
263	The role of free-living protozoa in protecting foodborne pathogens. , 2015, , 81-101.		1
264	Metatranscriptomic census of active protists in soils. ISME Journal, 2015, 9, 2178-2190.	4.4	274
265	Lateral gene transfers and the origins of the eukaryote proteome: a view from microbial parasites. Current Opinion in Microbiology, 2015, 23, 155-162.	2.3	42
266	Marked seasonality and high spatial variability of protist communities in shallow freshwater systems. ISME Journal, 2015, 9, 1941-1953.	4.4	165
267	Ciliary heterogeneity within a single cell: The Paramecium model. Methods in Cell Biology, 2015, 127, 457-485.	0.5	14
268	Cancer across the tree of life: cooperation and cheating in multicellularity. Philosophical Transactions of the Royal Society B: Biological Sciences, 2015, 370, 20140219.	1.8	303
269	Suigetsumonas clinomigrationis gen. et sp. nov., a Novel Facultative Anaerobic Nanoflagellate Isolated from the Meromictic Lake Suigetsu, Japan. Protist, 2015, 166, 409-421.	0.6	11
270	Root of Dictyostelia based on 213 universal proteins. Molecular Phylogenetics and Evolution, 2015, 92, 53-62.	1.2	16
271	Protistan Diversity in Environmental Molecular Surveys. , 2015, , 3-21.		12
272	Oldest Fossil Records of Marine Protists and the Geologic History Toward the Establishment of the Modern-Type Marine Protist World. , 2015, , 359-394.		8
273	Phaeodaria: Diverse Marine Cercozoans of World-Wide Distribution. , 2015, , 223-249.		25
274	Biology of Parasitic Heterotrophic Nanoflagellates: Parasitoids of Diatoms. , 2015, , 519-530.		1
275	Use of functional diversity to assess determinant assembly processes of testate amoebae community. Aquatic Ecology, 2015, 49, 561-571.	0.7	19
276	Biology, Diversity and Ecology of Free-Living Heterotrophic Flagellates. , 2015, , 63-87.		1
277	Open Questions on the Origin of Eukaryotes. Trends in Ecology and Evolution, 2015, 30, 697-708.	4.2	107
278	Discovery of PPi-type Phosphoenolpyruvate Carboxykinase Genes in Eukaryotes and Bacteria. Journal of Biological Chemistry, 2015, 290, 23960-23970.	1.6	33

		CITATION REPORT		
#	Article		IF	CITATIONS
279	Endosymbiotic origin and differential loss of eukaryotic genes. Nature, 2015, 524, 427	⁷ -432.	13.7	251
280	Comparative Genomics of a Bacterivorous Green Alga Reveals Evolutionary Causalities Consequences of Phago-Mixotrophic Mode of Nutrition. Genome Biology and Evolutio 3047-3061.	and n, 2015, 7,	1.1	36
281	KLF/SP Transcription Factor Family Evolution: Expansion, Diversification, and Innovatio Eukaryotes. Genome Biology and Evolution, 2015, 7, 2289-2309.	n in	1.1	93
282	Morphology, ontogenesis and molecular characterization of Atractos contortus VörÃ Stichotricha aculeata Wrzesniowskiego, 1866 (Ciliophora, Stichotrichida) with conside their systematic positions. European Journal of Protistology, 2015, 51, 351-373.	¶svÃjry, 1950 and eration of	0.5	14
283	Multiple origins of Heliozoa from flagellate ancestors: New cryptist subphylum Corbine superclass Corbistoma, and monophyly of Haptista, Cryptista, Hacrobia and Chromista Phylogenetics and Evolution, 2015, 93, 331-362.		1.2	73
284	Evolution of Proteasome Regulators in Eukaryotes. Genome Biology and Evolution, 20	15, 7, 1363-1379.	1.1	77
285	Metabolic Capacity of Mitochondrion-related Organelles in the Free-living Anaerobic St Cantina marsupialis. Protist, 2015, 166, 534-550.	tramenopile	0.6	12
286	Sex or no sex? Group I introns and independent marker genes reveal the existence of t reproductively isolated biospecies in Trichia varia (Myxomycetes). Organisms Diversity 2015, 15, 631-650.	hree sexual but and Evolution,	0.7	43
287	The TOM Complex of Amoebozoans: the Cases of the Amoeba Acanthamoeba castella Mold Dictyostelium discoideum. Protist, 2015, 166, 349-362.	nii and the Slime	0.6	15
288	The position of "lower plants―on the tree of life. Biology Bulletin, 2015, 42, 500-	507.	0.1	2
289	Flagellar motility in eukaryotic human parasites. Seminars in Cell and Developmental B 113-127.	iiology, 2015, 46,	2.3	35
290	Calcium signalling in the ciliated protozoan model, Paramecium: Strict signal localisati epigenetically controlled positioning of different Ca2+-channels. Cell Calcium, 2015, 5		1.1	20
291	Agglutinated versus microgranular foraminifers: end of a paradigm?. Journal of System Palaeontology, 2015, 13, 75-95.	atic	0.6	28
292	Experimental evidence for strong stabilizing forces at high functional diversity of aqua communities. Ecology, 2015, 96, 1340-1350.	tic microbial	1.5	40
293	Big answers from small worlds: a user's guide for protist microcosms as a model system and evolution. Methods in Ecology and Evolution, 2015, 6, 218-231.	m in ecology	2.2	157
294	Protistan diversity in a permanently stratified meromictic lake (Lake <scp>A</scp> lats	ee, <scp>SW) Tj ETQq1 1 (</scp>	0.784314 1.8	rgBJ /Overlo
295	Ethylene in Plants. , 2015, , .			28
296	Phylogeny and Reclassification of <i>Hemistasia phaeocysticola</i> (Scherffel) ElbrÃæ Schnepf, 1996. Journal of Eukaryotic Microbiology, 2015, 62, 426-429.	nter &	0.8	39

#	Article	IF	CITATIONS
297	Strains of the Morphospecies <i>Ploeotia costata</i> (Euglenozoa) Isolated from the Western North Pacific (Taiwan) Reveal Substantial Genetic Differences. Journal of Eukaryotic Microbiology, 2015, 62, 318-326.	0.8	6
298	Distribution of small phytoflagellates along an <scp>A</scp> rctic fjord transect. Environmental Microbiology, 2015, 17, 2393-2406.	1.8	33
299	Impact of long-term fertilizer treatment on the microeukaryotic community structure of a rice field soil. Soil Biology and Biochemistry, 2015, 80, 237-243.	4.2	40
300	Multigene phylogeny resolves deep branching of Amoebozoa. Molecular Phylogenetics and Evolution, 2015, 83, 293-304.	1.2	84
301	Complex communities of small protists and unexpected occurrence of typical marine lineages in shallow freshwater systems. Environmental Microbiology, 2015, 17, 3610-3627.	1.8	80
302	Methodological Studies on Estimates of Abundance and Diversity of Heterotrophic Flagellates from the Deep-Sea Floor. Journal of Marine Science and Engineering, 2016, 4, 22.	1.2	11
303	Ciliate Pheromones. , 2016, , 1-16.		2
304	Proximity-Dependent Biotin Identification (BioID) in Dictyostelium Amoebae. Methods in Enzymology, 2016, 569, 23-42.	0.4	29
305	Lipidomic Approaches towards Deciphering Glycolipids from Microalgae as a Reservoir of Bioactive Lipids. Marine Drugs, 2016, 14, 101.	2.2	96
306	Phylogeny and Classification of Euglenophyceae: A Brief Review. Frontiers in Ecology and Evolution, 2016, 4, .	1.1	15
307	Identification of Plasmodium falciparum Translation Initiation eIF2Î ² Subunit: Direct Interaction with Protein Phosphatase Type 1. Frontiers in Microbiology, 2016, 7, 777.	1.5	18
308	Resilience of Freshwater Communities of Small Microbial Eukaryotes Undergoing Severe Drought Events. Frontiers in Microbiology, 2016, 7, 812.	1.5	26
309	Widespread Inter- and Intra-Domain Horizontal Gene Transfer of d-Amino Acid Metabolism Enzymes in Eukaryotes. Frontiers in Microbiology, 2016, 7, 2001.	1.5	28
310	An Evolutionary Framework for Understanding the Origin of Eukaryotes. Biology, 2016, 5, 18.	1.3	23
311	The Diversity of Ribonuclease P: Protein and RNA Catalysts with Analogous Biological Functions. Biomolecules, 2016, 6, 27.	1.8	62
312	The Trypanosome Flagellar Pocket Collar and Its Ring Forming Protein—TbBILBO1. Cells, 2016, 5, 9.	1.8	19
313	Arginine deiminase pathway enzymes: evolutionary history in metamonads and other eukaryotes. BMC Evolutionary Biology, 2016, 16, 197.	3.2	40
314	On the reversibility of parasitism: adaptation to a free-living lifestyle via gene acquisitions in the diplomonad Trepomonas sp. PC1. BMC Biology, 2016, 14, 62.	1.7	38

#	Article	IF	CITATIONS
315	Analysis of the Microprocessor in Dictyostelium: The Role of RbdB, a dsRNA Binding Protein. PLoS Genetics, 2016, 12, e1006057.	1.5	16
316	Ultrastructural and Molecular Characterisation of an Heterosporis-Like Microsporidian in Australian Sea Snakes (Hydrophiinae). PLoS ONE, 2016, 11, e0150724.	1.1	8
317	Nosema ceranae Infection Promotes Proliferation of Yeasts in Honey Bee Intestines. PLoS ONE, 2016, 11, e0164477.	1.1	31
318	Dual RNA-Sequencing of Eucalyptus nitens during Phytophthora cinnamomi Challenge Reveals Pathogen and Host Factors Influencing Compatibility. Frontiers in Plant Science, 2016, 7, 191.	1.7	54
319	Next-Generation Sequencing — An Overview of the History, Tools, and "Omic―Applications. , 0, , .		94
320	Interactome Mapping Reveals the Evolutionary History of the Nuclear Pore Complex. PLoS Biology, 2016, 14, e1002365.	2.6	90
321	Comparative Metabolism of Freeâ€living <i>Bodo saltans</i> and Parasitic Trypanosomatids. Journal of Eukaryotic Microbiology, 2016, 63, 657-678.	0.8	86
322	The evolution of silicon transporters in diatoms. Journal of Phycology, 2016, 52, 716-731.	1.0	44
323	Morphological and molecular characterization of <i>Ptychodiscus noctiluca</i> revealed the polyphyletic nature of the order Ptychodiscales (Dinophyceae). Journal of Phycology, 2016, 52, 793-805.	1.0	6
324	<i>Phytophthora infestans</i> Argonaute 1 binds micro <scp>RNA</scp> and small <scp>RNA</scp> s from effector genes and transposable elements. New Phytologist, 2016, 211, 993-1007.	3.5	41
325	Comparative Cell Biology and Evolution of Annexins in Diplomonads. MSphere, 2016, 1, .	1.3	9
326	Morphology, Ontogeny, and Molecular Phylogeny of Two Freshwater Species of <i>Deviata</i> (Ciliophora, Hypotrichia) from Southern China. Journal of Eukaryotic Microbiology, 2016, 63, 771-785.	0.8	16
327	Species- and Strain-Specific Adaptation of the HSP70 Super Family in Pathogenic Trypanosomatids. Genome Biology and Evolution, 2016, 8, 1980-1995.	1.1	20
328	<i>Nematopsis temporariae</i> (Gregarinasina, Apicomplexa, Alveolata) is an intracellular infectious agent of tadpole livers. Environmental Microbiology Reports, 2016, 8, 675-679.	1.0	18
329	Energy for two: New archaeal lineages and the origin of mitochondria. BioEssays, 2016, 38, 850-856.	1.2	31
330	Thorough highâ€ŧhroughput sequencing analyses unravels huge diversities of soil parasitic protists. Environmental Microbiology, 2016, 18, 1669-1672.	1.8	36
331	<i>>Vermistella arctica</i> n. sp. Nominates the Genus <i>>Vermistella</i> as a Candidate for Taxon with Bipolar Distribution. Journal of Eukaryotic Microbiology, 2016, 63, 210-219.	0.8	4
332	Microbial eukaryotic distributions and diversity patterns in a deepâ€sea methane seep ecosystem. Environmental Microbiology, 2016, 18, 3022-3043.	1.8	40

#	Article	IF	CITATIONS
333	The Morphology, Ultrastructure and SSU rRNA Gene Sequence of a New Freshwater Flagellate, <i>Neobodo borokensis</i> n. sp. (Kinetoplastea, Excavata). Journal of Eukaryotic Microbiology, 2016, 63, 220-232.	0.8	26
334	<i>Sorodiplophrys stercorea</i> : Another Novel Lineage of Sorocarpic Multicellularity. Journal of Eukaryotic Microbiology, 2016, 63, 623-628.	0.8	28
335	Nextâ€Generation Sequencing Assessment of Eukaryotic Diversity in Oil Sands Tailings Ponds Sediments and SurfaceÂWater. Journal of Eukaryotic Microbiology, 2016, 63, 732-743.	0.8	26
336	Heme pathway evolution in kinetoplastid protists. BMC Evolutionary Biology, 2016, 16, 109.	3.2	19
337	The All-Data-Based Evolutionary Hypothesis of Ciliated Protists with a Revised Classification of the Phylum Ciliophora (Eukaryota, Alveolata). Scientific Reports, 2016, 6, 24874.	1.6	271
338	The unconventional kinetoplastid kinetochore: from discovery toward functional understanding. Biochemical Society Transactions, 2016, 44, 1201-1217.	1.6	20
339	Plant Proteins Are Smaller Because They Are Encoded by Fewer Exons than Animal Proteins. Genomics, Proteomics and Bioinformatics, 2016, 14, 357-370.	3.0	43
340	<i>Blastocystis</i> Mitochondrial Genomes Appear to Show Multiple Independent Gains and Losses of Start and Stop Codons. Genome Biology and Evolution, 2016, 8, 3340-3350.	1.1	9
341	A calcium sensor – protein kinase signaling module diversified in plants and is retained in all lineages of Bikonta species. Scientific Reports, 2016, 6, 31645.	1.6	34
343	Protist systematics, ecology and next generation sequencing. , 0, , 195-216.		0
344	Expansion of the molecular and morphological diversity of Acanthamoebidae (Centramoebida,) Tj ETQq0 0 0 rgBT	Overlock	2 10 Tf 50 34
345	Animal–Symbiodinium Symbioses: Foundations of Coral Reef Ecosystems. Advances in Environmental Microbiology, 2016, , 269-294.	0.1	16
346	Modern benthic foraminiferal diversity of Jeju Island and initial insights into the total foraminiferal diversity of Korea. Marine Biodiversity, 2016, 46, 337-354.	0.3	15
347	Diversity of extracellular proteins during the transition from the â€~proto-apicomplexan' alveolates to the apicomplexan obligate parasites. Parasitology, 2016, 143, 1-17.	0.7	56
348	Complex evolution of two types of cardiolipin synthase in the eukaryotic lineage stramenopiles. Molecular Phylogenetics and Evolution, 2016, 101, 133-141.	1.2	22
349	Losing Complexity: The Role of Simplification in Macroevolution. Trends in Ecology and Evolution, 2016, 31, 608-621.	4.2	55
350	Fungal diversity notes 253–366: taxonomic and phylogenetic contributions to fungal taxa. Fungal Diversity, 2016, 78, 1-237.	4.7	239
351	Bacterial and eukaryal diversity in soils forming from acid mine drainage precipitates under reclaimed vegetation and biological crusts. Applied Soil Ecology, 2016, 105, 57-66.	2.1	32

#	Article	IF	CITATIONS
352	Taxonomy and molecular systematics of three oligotrich (s.l.) ciliates including descriptions of two new species, <i>Strombidium guangdongense</i> sp. nov. and <i>Strombidinopsis sinicum</i> sp. nov. (Protozoa, Ciliophora). Systematics and Biodiversity, 2016, 14, 452-465.	0.5	20
353	Regulation of nuclear shape and size in plants. Current Opinion in Cell Biology, 2016, 40, 114-123.	2.6	25
354	Uniquely designed nuclear structures of lower eukaryotes. Current Opinion in Cell Biology, 2016, 40, 66-73.	2.6	9
355	Investigating the diversity of the 18S SSU rRNA hyper-variable region of Theileria in cattle and Cape buffalo (Syncerus caffer) from southern Africa using a next generation sequencing approach. Ticks and Tick-borne Diseases, 2016, 7, 869-879.	1.1	12
356	A small mitochondrial protein present in myzozoans is essential for malaria transmission. Open Biology, 2016, 6, 160034.	1.5	17
357	New kid on the block – the clubroot pathogen genome moves the plasmodiophorids into the genomic era. European Journal of Plant Pathology, 2016, 145, 531-542.	0.8	30
358	Reducing long-branch effects in multi-protein data uncovers a close relationship between Alveolata and Rhizaria. Molecular Phylogenetics and Evolution, 2016, 101, 1-7.	1.2	25
359	Occurrence and Activity of Slime Nets,Labyrinthulasp. Among Aquatic Plants in Cold and Oligohaline Baltic Sea Waters. Annales Botanici Fennici, 2016, 53, 139-143.	0.0	5
360	A Eukaryote without a Mitochondrial Organelle. Current Biology, 2016, 26, 1274-1284.	1.8	302
361	Apoptosis: its origin, history, maintenance and the medical implications for cancer and aging. Physical Biology, 2016, 13, 031001.	0.8	177
362	A Novel Lineage of â€~Naked Filose Amoebae'; Kraken carinae gen. nov. sp. nov. (Cercozoa) with a Remarkable Locomotion by Disassembly of its Cell Body. Protist, 2016, 167, 268-278.	0.6	19
363	Multigene-based phylogeny of the ciliate families Amphisiellidae and Trachelostylidae (Protozoa:) Tj ETQq1 🕻	1 0.784314 rgBT 1.2	/Qverlock]
364	Local and intercontinental comparisons of test morphology in the little-known testate amoeba Cyphoderia laevis Penard. European Journal of Protistology, 2016, 56, 67-78.	0.5	0
365	LINCing the eukaryotic tree of life – towards a broad evolutionary comparison of nucleocytoplasmic bridging complexes. Journal of Cell Science, 2016, 129, 3523-3531.	1.2	37
366	Genome microsatellite diversity within the Apicomplexa phylum. Molecular Genetics and Genomics, 2016, 291, 2117-2129.	1.0	1
367	Oligotrophic lagoons of the <scp>S</scp> outh <scp>P</scp> acific Ocean are home to a surprising number of novel eukaryotic microorganisms. Environmental Microbiology, 2016, 18, 4549-4563.	1.8	23
368	The ultrastructure of the zoospores of the parasitic dinoflagellate Ichthyodinium chabelardi Hollande et J. Cachon, 1952 (Alveolata: Dinoflagellata). Doklady Biological Sciences, 2016, 468, 125-128.	0.2	3
369	Diversification rates and species richness across the Tree of Life. Proceedings of the Royal Society B: Biological Sciences, 2016, 283, 20161334.	1.2	113

#	ARTICLE	IF	CITATIONS
370	Phylogenetic Resolution of Deep Eukaryotic and Fungal Relationships Using Highly Conserved Low-Copy Nuclear Genes. Genome Biology and Evolution, 2016, 8, 2683-2701.	1.1	31
371	Evolution in the Cycles of Life. Annual Review of Genetics, 2016, 50, 133-154.	3.2	99
372	Cinderella's helping pigeons of the microbial world: The potential of testate amoebae for identifying cryptotephra. European Journal of Protistology, 2016, 55, 152-164.	0.5	10
373	Exploiting the Achilles' heel of membrane trafficking in trypanosomes. Current Opinion in Microbiology, 2016, 34, 97-103.	2.3	28
374	Fatty acid production of tropical thraustochytrids from Malaysian mangroves. Botanica Marina, 2016, 59, 321-338.	0.6	17
375	The changing view of eukaryogenesis – fossils, cells, lineages and how they all come together. Journal of Cell Science, 2016, 129, 3695-3703.	1.2	77
376	The Evolution of Silicon Transport in Eukaryotes. Molecular Biology and Evolution, 2016, 33, 3226-3248.	3.5	107
377	Fine-Tuning Motile Cilia and Flagella: Evolution of the Dynein Motor Proteins from Plants to Humans at High Resolution. Molecular Biology and Evolution, 2016, 33, 3249-3267.	3.5	54
378	Systematics of organic-walled microfossils from the ca. 780–740 Ma Chuar Group, Grand Canyon, Arizona. Journal of Paleontology, 2016, 90, 815-853.	0.5	51
379	Terpene synthase genes in eukaryotes beyond plants and fungi: Occurrence in social amoebae. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 12132-12137.	3.3	92
380	Ciliate community structure and interactions within the planktonic food web in two alpine lakes of contrasting transparency. Freshwater Biology, 2016, 61, 1950-1965.	1.2	22
381	A Phylogenomic Framework to Study the Diversity and Evolution of Stramenopiles (=Heterokonts). Molecular Biology and Evolution, 2016, 33, 2890-2898.	3.5	125
382	Trypanosome <scp>RNA</scp> editing: the complexity of getting U in and taking U out. Wiley Interdisciplinary Reviews RNA, 2016, 7, 33-51.	3.2	124
383	Absence of a conventional spindle mitotic checkpoint in the binucleated single-celled parasite Giardia intestinalis. European Journal of Cell Biology, 2016, 95, 355-367.	1.6	17
384	Repertory of eukaryotes (eukaryome) in the human gastrointestinal tract: taxonomy and detection methods. Parasite Immunology, 2016, 38, 12-36.	0.7	47
385	The nonopisthokont septins. Methods in Cell Biology, 2016, 136, 1-19.	0.5	10
386	An Estimation of the Global Diversity and Distribution of the Smallest Eukaryotes: Biogeography of Marine Benthic Heterotrophic Flagellates. Protist, 2016, 167, 411-424.	0.6	31
387	Light- and Electron-microscopical Study of Belonocystis marina sp. nov. (Eukaryota: incertae sedis). Protist, 2016, 167, 479-489.	0.6	1

		EPORT	
#	Article	IF	CITATIONS
388	Multiciliated Cells in Animals. Cold Spring Harbor Perspectives in Biology, 2016, 8, a028233.	2.3	91
390	Placement of the unclassified Cyranomonas australis Lee 2002 within a novel clade of Cercozoa. European Journal of Protistology, 2016, 56, 60-66.	0.5	2
391	From writing to reading the encyclopedia of life. Philosophical Transactions of the Royal Society B: Biological Sciences, 2016, 371, 20150321.	1.8	48
392	Extreme Diversity of Diplonemid Eukaryotes in the Ocean. Current Biology, 2016, 26, 3060-3065.	1.8	105
393	Integrated omics study of lipid droplets from Plasmodiophora brassicae. Scientific Reports, 2016, 6, 36965.	1.6	59
394	MyTH4-FERM myosins have an ancient and conserved role in filopod formation. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, E8059-E8068.	3.3	24
395	Unikonts, Evolution and Diversification of (with Emphasis on Fungal-Like Forms). , 2016, , 325-332.		5
396	Higher classification and phylogeny of Euglenozoa. European Journal of Protistology, 2016, 56, 250-276.	0.5	63
397	What do isogamous organisms teach us about sex and the two sexes?. Philosophical Transactions of the Royal Society B: Biological Sciences, 2016, 371, 20150532.	1.8	46
398	It's official – Cryptosporidium is a gregarine: What are the implications for the water industry?. Water Research, 2016, 105, 305-313.	5.3	110
399	The protein domains of the Dictyostelium microprocessor that are required for correct subcellular localization and for microRNA maturation. RNA Biology, 2016, 13, 1000-1010.	1.5	12
400	The bacterial-fungal energy channel concept challenged by enormous functional versatility of soil protists. Soil Biology and Biochemistry, 2016, 102, 22-25.	4.2	129
401	Delineation of six species of the primitive algal genus Glaucocystis based on in situ ultrastructural characteristics. Scientific Reports, 2016, 6, 29209.	1.6	13
402	Protist Diversification. , 2016, , 344-360.		13
403	Genotype diversity in the honey bee parasite Nosema ceranae: multi-strain isolates, cryptic sex or both?. BMC Evolutionary Biology, 2016, 16, 216.	3.2	11
404	Phylogeny and Evolution. , 2016, , 383-408.		4
405	Planktonic protistan communities in lakes along a large-scale environmental gradient. FEMS Microbiology Ecology, 2017, 93, fiw231.	1.3	28
406	Mitochondrial clock: moderating evolution of early eukaryotes in light of the Proterozoic oceans. Biologia (Poland), 2016, 71, 843-852.	0.8	0

#	Article	IF	Citations
407	Biogeography of Radiolaria Polycystina (Protista) in the World Ocean. Progress in Oceanography, 2016, 149, 82-105.	1.5	44
408	Comparative Ultrastructure of Fornicate Excavates, Including a Novel Free-living Relative of Diplomonads: Aduncisulcus paluster gen. et sp. nov Protist, 2016, 167, 584-596.	0.6	20
409	Jakobida. , 2016, , 1-32.		1
410	Preaxostyla. , 2016, , 1-36.		3
411	Triassic chrysophyte cyst fossils discovered in the Ordos Basin, China. Geology, 2016, 44, 1031-1034.	2.0	26
412	Yeast Ivy1p Is a Putative I-BAR-domain Protein with pH-sensitive Filament Forming Ability <i>in vitro</i> . Cell Structure and Function, 2016, 41, 1-11.	0.5	12
413	Morphology, Cell Division, and Phylogeny of <i>Uroleptus longicaudatus</i> (Ciliophora,) Tj ETQq0 0 0 rgBT /Ove 2016, 63, 349-362.	rlock 10 T 0.8	f 50 507 Td 16
414	<i>Trachyrhizium urniformis</i> n. g., n. sp., a Novel Marine Filose Thecate Amoeba Related to a Cercozoan Environmental Clade (Novel Clade 4). Journal of Eukaryotic Microbiology, 2016, 63, 722-731.	0.8	13
415	Phenology of cryptomonads and the CRY1 lineage in a coastal brackish lagoon (Vistula Lagoon, Baltic) Tj ETQqO	0 0 rgBT /0 1.0	Dvgglock 10 T
416	A paneukaryotic genomic analysis of the small GTPase RABL2 underscores the significance of recurrent gene loss in eukaryote evolution. Biology Direct, 2016, 11, 5.	1.9	22
417	Highly efficient lipid production in the green alga Parachlorella kessleri: draft genome and transcriptome endorsed by whole-cell 3D ultrastructure. Biotechnology for Biofuels, 2016, 9, 13.	6.2	56
418	Characterization of recombinase DMC1B and its functional role as Rad51 in DNA damage repair in Giardia duodenalis trophozoites. Biochimie, 2016, 127, 173-186.	1.3	7
419	AGGLUTINATED OR PORCELANEOUS TESTS: WHERE TO DRAW THE LINE?. Journal of Foraminiferal Research, 2016, 46, 333-344.	0.1	15
420	Morphology and Molecular Phylogeny of Coelomic Gregarines (Apicomplexa) with Different Types of Motility: Urospora ovalis and U. travisiae from the Polychaete Travisia forbesii. Protist, 2016, 167, 279-301.	0.6	10
421	Current status of Blastocystis: A personal view. Parasitology International, 2016, 65, 763-771.	0.6	253
422	Agglutination of benthic foraminifera in relation to mesoscale bathymetric features in the abyssal NE Atlantic (Porcupine Abyssal Plain). Marine Micropaleontology, 2016, 123, 15-28.	0.5	15
423	What an Intron May Tell: Several Sexual Biospecies Coexist in Meriderma spp. (Myxomycetes). Protist, 2016, 167, 234-253.	0.6	33
424	Intermediate introns in nuclear genes of euglenids – are they a distinct type?. BMC Evolutionary Biology, 2016, 16, 49.	3.2	15

#	Article	IF	CITATIONS
425	Protein import complexes in the mitochondrial outer membrane of Amoebozoa representatives. BMC Genomics, 2016, 17, 99.	1.2	6
426	The Species Problem in Myxomycetes Revisited. Protist, 2016, 167, 319-338.	0.6	30
427	Cutaneous myxidiosis in European eel, <i>Anguilla anguilla</i> (Linnaeus, 1758): histopathology, histochemistry and laminin immunohistochemistry. Journal of Fish Diseases, 2016, 39, 845-851.	0.9	2
428	A new type of 3â€D peripheral ultrastructure in <i>Glaucocystis</i> (Glaucocystales, Glaucophyta) as revealed by ultraâ€high voltage electron microscopy. Journal of Phycology, 2016, 52, 486-490.	1.0	2
429	The evolution of substrate specificityâ€associated residues and Ca ²⁺ â€binding motifs in <scp>EF</scp> â€handâ€containing type <scp>II NAD</scp> (P)H dehydrogenases. Physiologia Plantarum, 2016, 157, 338-351.	2.6	8
430	Fine-structural Observations on Siliceous Scale Production and Shell Assembly in the Testate Amoeba Paulinella chromatophora. Protist, 2016, 167, 303-318.	0.6	11
431	The evolution of ERMIONE in mitochondrial biogenesis and lipid homeostasis: An evolutionary view from comparative cell biology. Biochimica Et Biophysica Acta - Molecular and Cell Biology of Lipids, 2016, 1861, 900-912.	1.2	49
432	The soil food web revisited: Diverse and widespread mycophagous soil protists. Soil Biology and Biochemistry, 2016, 94, 10-18.	4.2	175
433	Isolation of diverse amoebal grazers of freshwater cyanobacteria for the development of model systems to study predator–prey interactions. Algal Research, 2016, 13, 85-93.	2.4	22
434	Soil biota in a megadiverse country: Current knowledge and future research directions in South Africa. Pedobiologia, 2016, 59, 129-174.	0.5	45
435	Detection of Balamuthia mandrillaris DNA in the storage case of contact lenses in Germany. Parasitology Research, 2016, 115, 2111-2114.	0.6	9
436	Diverged composition and regulation of theTrypanosoma bruceiorigin recognition complex that mediates DNA replication initiation. Nucleic Acids Research, 2016, 44, 4763-4784.	6.5	31
437	Rhinosporidium seeberiNuclear Cycle Activities Using Confocal Microscopy. Journal of Parasitology, 2016, 102, 60-68.	0.3	2
438	Systematics, Taxonomy and Species Names: Do They Matter?. , 2016, , 655-681.		24
439	Rhamnolipids as environmentally friendly biopesticide against plant pathogen <i>Phytophthora sojae</i> . Environmental Progress and Sustainable Energy, 2016, 35, 169-173.	1.3	22
440	Silicification in the Microalgae. , 2016, , 289-300.		16
441	Phylogenomics of †Discosea': A new molecular phylogenetic perspective on Amoebozoa with flat body forms. Molecular Phylogenetics and Evolution, 2016, 99, 144-154.	1.2	38
442	A Comparative Analysis of Mitochondrial Genomes in Eustigmatophyte Algae. Genome Biology and Evolution, 2016, 8, 705-722.	1.1	33

#	Article	IF	CITATIONS
443	Reproductive cyst and operculum formation in the Cambrian–Ordovician galeate-plexus microfossils. Gff, 2016, 138, 278-294.	0.4	9
444	Current and future perspectives on the systematics, taxonomy and nomenclature of testate amoebae. European Journal of Protistology, 2016, 55, 105-117.	0.5	75
445	Peroxisomes in parasitic protists. Molecular and Biochemical Parasitology, 2016, 209, 35-45.	0.5	47
446	Aerobic mitochondria of parasitic protists: Diverse genomes and complex functions. Molecular and Biochemical Parasitology, 2016, 209, 46-57.	0.5	24
447	Protist diversity on a nature reserve in NW England—With particular reference to their role in soil biogenic silicon pools. Pedobiologia, 2016, 59, 51-59.	0.5	37
448	Plant evolution and terrestrialization during Palaeozoic times—The phylogenetic context. Review of Palaeobotany and Palynology, 2016, 227, 4-18.	0.8	60
449	Ancestral reconstruction of tick lineages. Ticks and Tick-borne Diseases, 2016, 7, 509-535.	1.1	62
450	Molecular Identification of Soil Eukaryotes and Focused Approaches Targeting Protist and Faunal Groups Using High-Throughput Metabarcoding. Methods in Molecular Biology, 2016, 1399, 125-140.	0.4	11
451	De novo transcriptome assembly of Perkinsus olseni trophozoite stimulated in vitro with Manila clam (Ruditapes philippinarum) plasma. Journal of Invertebrate Pathology, 2016, 135, 22-33.	1.5	14
452	A new species of small acritarch with a porous wall structure from the early Cambrian of Estonia and implications for the fossil record of eukaryotic picoplankton. Palynology, 2016, 40, 343-356.	0.7	6
453	Light and electron microscopic observations of the reproductive swarmer cells of nassellarian and spumellarian polycystines (Radiolaria). European Journal of Protistology, 2016, 54, 19-32.	0.5	10
454	Dinoflagellate cyst production over an annual cycle in seasonally ice-covered Hudson Bay. Marine Micropaleontology, 2016, 125, 1-24.	0.5	45
455	Novel insights on ENTH domain-containing proteins in apicomplexan parasites. Parasitology Research, 2016, 115, 2191-2202.	0.6	3
456	Initiation of a reversal to uniseriality in the polymorphinid Nodocantabricus duplexmurus n. gen., n. sp.: A double-layered Foraminifera from the lower-middle Cenomanian of Cantabria, N-Spain. Cretaceous Research, 2016, 63, 14-22.	0.6	1
457	Testate Amoebae Like It Hot: Species Richness Decreases Along a Subalpine-Alpine Altitudinal Gradient in Both Natural Calluna vulgaris Litter and Transplanted Minuartia sedoides Cushions. Microbial Ecology, 2016, 71, 725-734.	1.4	14
458	Bacterial and eukaryotic biodiversity patterns in terrestrial and aquatic habitats in the SÃr Rondane Mountains, Dronning Maud Land, East Antarctica. FEMS Microbiology Ecology, 2016, 92, fiw041.	1.3	30
459	Group II Intron-Mediated <i>Trans</i> -Splicing in the Gene-Rich Mitochondrial Genome of an Enigmatic Eukaryote, <i>Diphylleia rotans</i> . Genome Biology and Evolution, 2016, 8, 458-466.	1.1	28
460	Protistan diversity and activity inferred from RNA and DNA at a coastal ocean site in the eastern North Pacific. FEMS Microbiology Ecology, 2016, 92, fiw050.	1.3	74

#	Article	IF	CITATIONS
461	New lineage of Triassic aragonitic Foraminifera and reassessment of the class Nodosariata. Journal of Systematic Palaeontology, 2016, 14, 919-938.	0.6	14
462	Legionella pneumophila, armed to the hilt: justifying the largest arsenal of effectors in the bacterial world. Current Opinion in Microbiology, 2016, 29, 74-80.	2.3	114
463	Actin evolution in ciliates (Protist, Alveolata) is characterized by high diversity and three duplication events. Molecular Phylogenetics and Evolution, 2016, 96, 45-54.	1.2	16
464	Abundance, diversity and community composition of free-living protozoa on vegetable sprouts. Food Microbiology, 2016, 55, 55-63.	2.1	13
465	The <i>Physarum polycephalum</i> Genome Reveals Extensive Use of Prokaryotic Two-Component and Metazoan-Type Tyrosine Kinase Signaling. Genome Biology and Evolution, 2016, 8, 109-125.	1.1	87
466	Zoosporic parasites infecting marine diatoms – A black box that needs to be opened. Fungal Ecology, 2016, 19, 59-76.	0.7	109
467	Multiple Origins of Eukaryotic <i>cox15</i> Suggest Horizontal Gene Transfer from Bacteria to Jakobid Mitochondrial DNA. Molecular Biology and Evolution, 2016, 33, 122-133.	3.5	21
468	Testate amoeba <i>Rhogostoma minus</i> Belar, 1921, associated with nodular gill disease of rainbow trout, <i>Oncorhynchus mykiss</i> (Walbaum). Journal of Fish Diseases, 2016, 39, 539-546.	0.9	20
469	Back to monoxeny: Phytomonas nordicus descended from dixenous plant parasites. European Journal of Protistology, 2016, 52, 1-10.	0.5	19
470	Ecological importance of soil bacterivores for ecosystem functions. Plant and Soil, 2016, 398, 1-24.	1.8	251
471	Algal affinities of Ediacaran and Cambrian organic-walled microfossils with internal reproductive bodies: <i>Tanarium</i> and other morphotypes. Palynology, 2016, 40, 83-121.	0.7	32
472	Molecular organisms. Biology and Philosophy, 2016, 31, 571-589.	0.7	0
473	Repertoire of human gut microbes. Microbial Pathogenesis, 2017, 106, 103-112.	1.3	70
474	Dinoflagellate taxonomy — a review and proposal of a revised classification. Marine Biodiversity, 2017, 47, 381-403.	0.3	46
475	Lipid quantification techniques for screening oleaginous species of microalgae for biofuel production. European Journal of Lipid Science and Technology, 2017, 119, 1500469.	1.0	17
476	Dietaryâ€driven variation effects on the symbiotic flagellate protist communities of the subterranean termite <i>Reticulitermes grassei</i> Clément. Journal of Applied Entomology, 2017, 141, 300-307.	0.8	14
477	<i>Neovahlkampfia nana</i> n. sp. Reinforcing an Underrepresented Subclade of Tetramitia, Heterolobosea. Journal of Eukaryotic Microbiology, 2017, 64, 78-87.	0.8	8
478	Signalling in ciliates: long- and short-range signals and molecular determinants for cellular dynamics. Biological Reviews, 2017, 92, 60-107.	4.7	25

#	Article	IF	CITATIONS
479	Loss of GET pathway orthologs in <i>Arabidopsis thaliana</i> causes root hair growth defects and affects SNARE abundance. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E1544-E1553.	3.3	56
480	Beyond the "Code― A Guide to the Description and Documentation of Biodiversity in Ciliated Protists (Alveolata, Ciliophora). Journal of Eukaryotic Microbiology, 2017, 64, 539-554.	0.8	108
481	<scp>DNA</scp> metabarcoding reveals that 200â€ <i>μ</i> mâ€sizeâ€fractionated filtering is unable to discriminate between planktonic microbial and large eukaryotes. Molecular Ecology Resources, 2017, 17, 991-1002.	2.2	50
482	Cytokinesis in Metazoa and Fungi. Cold Spring Harbor Perspectives in Biology, 2017, 9, a022343.	2.3	63
483	Polymetallic nodules, sediments, and deep waters in the equatorial North Pacific exhibit highly diverse and distinct bacterial, archaeal, and microeukaryotic communities. MicrobiologyOpen, 2017, 6, e00428.	1.2	93
484	Amoebozoans are Secretly but Ancestrally Sexual: Evidence for Sex Genes and Potential Novel Crossover Pathways in Diverse Groups of Amoebae. Genome Biology and Evolution, 2017, 9, evx002.	1.1	28
485	Pathogens of brown algae: culture studies of <i>Anisolpidium ectocarpii</i> and <i>A. rosenvingei</i> reveal that the Anisolpidiales are uniflagellated oomycetes. European Journal of Phycology, 2017, 52, 133-148.	0.9	34
486	Discovering Protein-Coding Genes from the Environment: Time for the Eukaryotes?. Trends in Biotechnology, 2017, 35, 824-835.	4.9	18
487	Lineage‧pecific and Highly Derived Gene Sequences Among Amoebozoa, Revealed by the Comparative Analysis of Transcriptomes from Twelve Amoebozoan Species. Journal of Eukaryotic Microbiology, 2017, 64, 622-631.	0.8	0
488	Molecular phylogeny of extant Holothuroidea (Echinodermata). Molecular Phylogenetics and Evolution, 2017, 111, 110-131.	1.2	133
489	Ultrastructure and molecular phylogenetic position of a new marine sand-dwelling dinoflagellate from British Columbia, Canada: <i>Pseudadenoides polypyrenoides</i> sp. nov. (Dinophyceae). European Journal of Phycology, 2017, 52, 208-224.	0.9	7
490	Diversity of the Photosynthetic Paulinella Species, with the Description of Paulinella micropora sp. nov. and the Chromatophore Genome Sequence for strain KR01. Protist, 2017, 168, 155-170.	0.6	28
491	Fungi: Characteristics and Classification. , 2017, , 1-15.		10
492	The Protistan Microbiome of Grassland Soil: Diversity in the Mesoscale. Protist, 2017, 168, 546-564.	0.6	36
493	Soil protistology rebooted: 30 fundamental questions to start with. Soil Biology and Biochemistry, 2017, 111, 94-103.	4.2	130
494	An updated classification of rotaliid foraminifera based on ribosomal DNA phylogeny. Marine Micropaleontology, 2017, 132, 18-34.	0.5	46
495	Evolutionary Origin of Euglena. Advances in Experimental Medicine and Biology, 2017, 979, 3-17.	0.8	35
496	Euglena gracilis Genome and Transcriptome: Organelles, Nuclear Genome Assembly Strategies and Initial Features. Advances in Experimental Medicine and Biology, 2017, 979, 125-140.	0.8	35

C	ΓΑΤ	 . n		<u></u>	-
			F D	าเว	
\sim					

#	Article	IF	CITATIONS
497	Molecular characterisation of protistan species and communities in ships' ballast water across three U.S. coasts. Diversity and Distributions, 2017, 23, 680-691.	1.9	17
498	Ultrastructural and immunocytochemical investigation of paramylon combined with new 18S rDNA-based secondary structure analysis clarifies phylogenetic affiliation of Entosiphon sulcatum (Euglenida: Euglenozoa). Organisms Diversity and Evolution, 2017, 17, 509-520.	0.7	4
499	Expansion of the redox-sensitive proteome coincides with the plastid endosymbiosis. Nature Plants, 2017, 3, 17066.	4.7	26
500	Between a Pod and a Hard Test: The Deep Evolution of Amoebae. Molecular Biology and Evolution, 2017, 34, 2258-2270.	3.5	161
501	A Comparison of Methods to Analyze Aquatic Heterotrophic Flagellates of Different Taxonomic Groups. Protist, 2017, 168, 375-391.	0.6	5
502	Systematic studies on ciliates (Alveolata, Ciliophora) in China: Progress and achievements based on molecular information. European Journal of Protistology, 2017, 61, 409-423.	0.5	64
503	Insights into Ciliary Genes and Evolution from Multi-Level Phylogenetic Profiling. Molecular Biology and Evolution, 2017, 34, 2016-2034.	3.5	54
504	Comparative interactomics provides evidence for functional specialization of the nuclear pore complex. Nucleus, 2017, 8, 340-352.	0.6	16
505	Further consideration on the phylogeny of the Ciliophora: Analyses using both mitochondrial and nuclear data with focus on the extremely confused class Phyllopharyngea. Molecular Phylogenetics and Evolution, 2017, 112, 96-106.	1.2	28
506	Diversity of organic-walled microfossils from the early Mesoproterozoic Ruyang Group, North China Craton – A window into the early eukaryote evolution. Precambrian Research, 2017, 297, 101-130.	1.2	84
507	Outerwear through the ages: evolutionary cell biology of vesicle coats. Current Opinion in Cell Biology, 2017, 47, 108-116.	2.6	56
508	Morphologic and phylogenetic studies of two hypotrichous ciliates, with notes on morphogenesis in Gastrostyla steinii Engelmann, 1862 (Ciliophora, Hypotrichia). European Journal of Protistology, 2017, 60, 119-133.	0.5	16
509	Identification of a new subtilisin-like protease NbSLP2 interacting with cytoskeletal protein septin in Microsporidia Nosema bombycis. Journal of Invertebrate Pathology, 2017, 148, 110-117.	1.5	7
510	Human Chitotriosidase: Catalytic Domain or Carbohydrate Binding Module, Who's Leading HCHT's Biological Function. Scientific Reports, 2017, 7, 2768.	1.6	14
511	Evolution: Of X-Cells and X-Men. Current Biology, 2017, 27, R408-R409.	1.8	0
512	An 18S rRNA Workflow for Characterizing Protists in Sewage, with a Focus on Zoonotic Trichomonads. Microbial Ecology, 2017, 74, 923-936.	1.4	36
513	Phylogenomic analysis of integral diiron membrane histidine motif-containing enzymes in ciliates provides insights into their function and evolutionary relationships. Molecular Phylogenetics and Evolution, 2017, 114, 1-13.	1.2	8
514	The Physiology of Phagocytosis in the Context of Mitochondrial Origin. Microbiology and Molecular Biology Reviews, 2017, 81, .	2.9	84

#	Article	IF	CITATIONS
515	Cyanobacterial Endosymbionts of Paulinella chromatophora Shed New Light on the Birth of Cellular Organelles. , 2017, , 279-328.		1
516	Symbiosis between Testate Amoebae and Photosynthetic Organisms. , 2017, , 399-419.		10
517	SILVA, RDP, Greengenes, NCBI and OTT — how do these taxonomies compare?. BMC Genomics, 2017, 18, 114.	1.2	327
518	Structures related to attachment and motility in the marine eugregarine Cephaloidophora cf. communis (Apicomplexa). European Journal of Protistology, 2017, 59, 1-13.	0.5	8
519	Key Ecological Roles for Zoosporic True Fungi in Aquatic Habitats. Microbiology Spectrum, 2017, 5, .	1.2	18
520	How Embryophytic is the Biosynthesis of Phenylpropanoids and their Derivatives in Streptophyte Algae?. Plant and Cell Physiology, 2017, 58, 934-945.	1.5	102
521	Changes in bacterial community composition and soil respiration indicate rapid successions of protist grazers during mineralization of maize crop residues. Pedobiologia, 2017, 62, 1-8.	0.5	37
522	<i>UniEuk</i> : Time to Speak a Common Language in Protistology!. Journal of Eukaryotic Microbiology, 2017, 64, 407-411.	0.8	74
523	Productivity gradient affects the temporal dynamics of testate amoebae in a neotropical floodplain. Ecological Indicators, 2017, 78, 264-269.	2.6	11
524	Effects of cyanobacteria Synechocystis spp. in the host-parasite model Crassostrea gasar–Perkinsus marinus. Aquatic Toxicology, 2017, 187, 100-107.	1.9	2
525	Evolutionary History of Subtilases in Land Plants and Their Involvement in Symbiotic Interactions. Molecular Plant-Microbe Interactions, 2017, 30, 489-501.	1.4	38
526	Origin and diversity of testate amoebae shell composition: Example of Bullinularia indica living in Sphagnum capillifolium. European Journal of Protistology, 2017, 59, 14-25.	0.5	3
527	Parasitism as the main factor shaping peptide vocabularies in current organisms. Parasitology, 2017, 144, 975-983.	0.7	7
528	Nuclear codon reassignments in the genomics era and mechanisms behind their evolution. BioEssays, 2017, 39, 1600221.	1.2	35
529	Parasites dominate hyperdiverse soil protist communities in Neotropical rainforests. Nature Ecology and Evolution, 2017, 1, 91.	3.4	262
530	Concerted Up-regulation of Aldehyde/Alcohol Dehydrogenase (ADHE) and Starch in Chlamydomonas reinhardtii Increases Survival under Dark Anoxia. Journal of Biological Chemistry, 2017, 292, 2395-2410.	1.6	26
531	Eukaryotic community diversity and spatial variation during drinking water production (by seawater) Tj ETQq0 0 Technology, 2017, 3, 92-105.	0 rgBT /Ov 1.2	verlock 10 Tf 9
532	SDH6 and SDH7 Contribute to Anchoring Succinate Dehydrogenase to the Inner Mitochondrial Membrane in <i>Arabidopsis thaliana</i> . Plant Physiology, 2017, 173, 1094-1108.	2.3	30

#	Article	IF	CITATIONS
533	The interaction of a Trypanosoma brucei KH-domain protein with a ribonuclease is implicated in ribosome processing. Molecular and Biochemical Parasitology, 2017, 211, 94-103.	0.5	11
534	Bothrosome Formation in Schizochytrium aggregatum (Labyrinthulomycetes, Stramenopiles) during Zoospore Settlement. Protist, 2017, 168, 206-219.	0.6	12
535	Relationships between Free-Living Amoeba and their Intracellular Bacteria. Proceedings of the Latvian Academy of Sciences, 2017, 71, 259-265.	0.0	3
536	Nine unanswered questions about cytokinesis. Journal of Cell Biology, 2017, 216, 3007-3016.	2.3	73
537	Deep-sea ciliates: Recorded diversity and experimental studies on pressure tolerance. Deep-Sea Research Part I: Oceanographic Research Papers, 2017, 128, 55-66.	0.6	16
538	Evolutionary Lessons from Species with Unique Kinetochores. Progress in Molecular and Subcellular Biology, 2017, 56, 111-138.	0.9	43
539	Prevalence of Pentatrichomonas hominis infections in six farmed wildlife species in Jilin, China. Veterinary Parasitology, 2017, 244, 160-163.	0.7	12
540	Haemoprotozoa: Making biological sense of molecular phylogenies. International Journal for Parasitology: Parasites and Wildlife, 2017, 6, 241-256.	0.6	47
541	Biotic interactions as drivers of algal origin and evolution. New Phytologist, 2017, 216, 670-681.	3.5	25
542	Morphology and phylogeny of the testate amoebae Euglypha bryophila Brown, 1911 and Euglypha cristata Leidy, 1874 (Rhizaria, Euglyphida). European Journal of Protistology, 2017, 61, 76-84.	0.5	4
543	Characterisation of sterol biosynthesis and validation of 14α-demethylase as a drug target in Acanthamoeba. Scientific Reports, 2017, 7, 8247.	1.6	38
544	First molecular characterization of Balantioides coli (Malmsten, 1857) isolates maintained in vitro culture and from feces of captive animals, Rio de Janeiro, Brazil. Veterinary Parasitology: Regional Studies and Reports, 2017, 10, 102-113.	0.3	9
545	Acidophilic green algal genome provides insights into adaptation to an acidic environment. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E8304-E8313.	3.3	93
546	A coarse-grained computational model of the nuclear pore complex predicts Phe-Gly nucleoporin dynamics. Journal of General Physiology, 2017, 149, 951-966.	0.9	8
547	Giardia's primitive GPL biosynthesis pathways with parasitic adaptation â€~patches': implications for Giardia's evolutionary history and for finding targets against Giardiasis. Scientific Reports, 2017, 7, 9507.	1.6	15
548	Centrin diversity and basal body patterning across evolution: new insights from Paramecium. Biology Open, 2017, 6, 765-776.	0.6	10
549	Intracellular metabolic pathway distribution in diatoms and tools for genome-enabled experimental diatom research. Philosophical Transactions of the Royal Society B: Biological Sciences, 2017, 372, 20160402.	1.8	38
550	The flagellar apparatus of the glaucophyte <i>Cyanophora cuspidata</i> . Journal of Phycology, 2017, 53, 1120-1150.	1.0	3

#	Article	IF	CITATIONS
551	Amoebozoan Lobose Amoebae (Tubulinea, Flabellinea, and Others). , 2017, , 1279-1309.		9
552	Protist Diversity and Eukaryote Phylogeny. , 2017, , 1-21.		35
553	Syllabus of Plant Families. A. Engler's Syllabus der Pflanzenfamilien, 13th edition. Part 2/1: Photoautotrophic Eukaryotic Algae. Glaucocystophyta, Cryptophyta, Dinophyta/Dinozoa, Haptophyta, Heterokontophyta/Ochrophyta, Chlorarachniophyta/ Cercozoa, Euglenophyta/Euglenozoa, Chlorophyta, Streptophyta p.p. W. Frey (ed.). Stuttgart: Borntraeger Science Publishers. 2015. 324 pp., 67 figures. ISBN 978 3 443 01083 6. 89 â,¬ (hardback) Edinburgh Journal of Botany, 2017, 74, 366-370.	0.4	0
554	Heterolobosea., 2017, , 1005-1046.		12
555	Protosteloid Amoebae (Protosteliida, Protosporangiida, Cavosteliida, Schizoplasmodiida,) Tj ETQq0 0 0 rgBT /Ove 1311-1348.	erlock 10	Tf 50 587 Td (6
556	Dictyostelia. , 2017, , 1433-1477.		3
557	Radiolaria and Phaeodaria. , 2017, , 731-763.		7
558	Apicomplexa. , 2017, , 567-624.		21
559	Paramyxida. , 2017, , 805-822.		1
560	Hyphochytriomycota and Oomycota. , 2017, , 435-505.		38
561	Centrohelida and Other Heliozoan-Like Protists. , 2017, , 955-971.		3
562	Retortamonadida (with Notes on Carpediemonas-Like Organisms and Caviomonadidae). , 2017, , 1247-1278.		4
563	Eustigmatophyceae. , 2017, , 367-406.		19
564	Zygnematophyta. , 2017, , 135-163.		1
565	Myxomycetes. , 2017, , 1405-1431.		9
566	Jakobida. , 2017, , 973-1003.		2
567	Preaxostyla. , 2017, , 1139-1174.		11
568	Parabasalia. , 2017, , 1175-1218.		8

#	Article	IF	CITATIONS
569	The Convoluted Evolution of Eukaryotes With Complex Plastids. Advances in Botanical Research, 2017, 84, 1-30.	0.5	20
570	Proposal of <scp><i>M</i></scp> <i>onorhizochytrium globosum</i> gen. nov., comb. nov. (<scp>S</scp> tramenopiles, <scp>L</scp> abyrinthulomycetes) for former <scp><i>T</i></scp> <i>hraustochytrium globosum</i> based on morphological features and phylogenetic relationships. Phycological Research. 2017. 65. 188-201.	0.8	17
571	Yeasts dominate soil fungal communities in three lowland Neotropical rainforests. Environmental Microbiology Reports, 2017, 9, 668-675.	1.0	14
572	Cryptomonads: A Model Organism Sheds Light on the Evolutionary History of Genome Reorganization in Secondary Endosymbioses. Advances in Botanical Research, 2017, 84, 263-320.	0.5	9
573	Anaeramoebidae fam. nov., a Novel Lineage of Anaerobic Amoebae and Amoeboflagellates of Uncertain Phylogenetic Position. Protist, 2017, 168, 495-526.	0.6	12
574	Application of Protists to Improve Plant Growth in Sustainable Agriculture. , 2017, , 263-273.		14
575	Are Thraustochytrids algae?. Fungal Biology, 2017, 121, 835-840.	1.1	84
576	Possible impacts of zoosporic parasites in diseases of commercially important marine mollusc species: part II. Labyrinthulomycota. Botanica Marina, 2017, 60, .	0.6	10
577	Functions of myosin motors tailored for parasitism. Current Opinion in Microbiology, 2017, 40, 113-122.	2.3	14
578	Let There Be Light: A Contemporary Primer on Primary Plastid Endosymbiosis. Advances in Botanical Research, 2017, 84, 31-56.	0.5	1
579	Secondary Plastids of Euglenophytes. Advances in Botanical Research, 2017, 84, 321-358.	0.5	8
580	Biology of Haptophytes: Complicated Cellular Processes Driving the Global Carbon Cycle. Advances in Botanical Research, 2017, 84, 219-261.	0.5	14
581	Antitrichomonal activity of δ opioid receptor antagonists, 7-benzylidenenaltrexone derivatives. Bioorganic and Medicinal Chemistry, 2017, 25, 4375-4383.	1.4	8
582	What Defines the "Kingdom―Fungi?. Microbiology Spectrum, 2017, 5, .	1.2	59
583	The calmodulin fused kinase novel gene family is the major system in plants converting Ca2+ signals to protein phosphorylation responses. Scientific Reports, 2017, 7, 4127.	1.6	12
584	Deciphering biodiversity and interactions between bacteria and microeukaryotes within epilithic biofilms from the Loue River, France. Scientific Reports, 2017, 7, 4344.	1.6	34
585	Heterotrophic flagellates of sphagnum bogs and lakes in Usman pine forest, Voronezh oblast. Inland Water Biology, 2017, 10, 182-191.	0.2	3
586	TCTE1 is a conserved component of the dynein regulatory complex and is required for motility and metabolism in mouse spermatozoa. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E5370-E5378.	3.3	74

#	Article	IF	CITATIONS
587	Analysis of triglyceride synthesis unveils a green algal soluble diacylglycerol acyltransferase and provides clues to potential enzymatic components of the chloroplast pathway. BMC Genomics, 2017, 18, 223.	1.2	25
588	Well-positioned nucleosomes punctuate polycistronic pol II transcription units and flank silent VSG gene arrays in Trypanosoma brucei. Epigenetics and Chromatin, 2017, 10, 14.	1.8	14
589	Longamoebia is not monophyletic: Phylogenomic and cytoskeleton analyses provide novel and well-resolved relationships of amoebozoan subclades. Molecular Phylogenetics and Evolution, 2017, 114, 249-260.	1.2	21
590	The family of berberine bridge enzyme-like enzymes: A treasure-trove of oxidative reactions. Archives of Biochemistry and Biophysics, 2017, 632, 88-103.	1.4	100
591	A comparative study of genome organization and epigenetic mechanisms in model ciliates, with an emphasis on Tetrahymena , Paramecium and Oxytricha. European Journal of Protistology, 2017, 61, 376-387.	0.5	33
592	Phylogeny mandalas for illustrating the Tree of Life. Molecular Phylogenetics and Evolution, 2017, 117, 168-178.	1.2	34
593	Probing the evolution, ecology and physiology of marine protists using transcriptomics. Nature Reviews Microbiology, 2017, 15, 6-20.	13.6	176
594	A six-gene phylogeny provides new insights into choanoflagellate evolution. Molecular Phylogenetics and Evolution, 2017, 107, 166-178.	1.2	59
595	Ultrastructure and Molecular Phylogeny of <i>lotanema spirale</i> gen. nov. et sp. nov., a New Lineage of Endobiotic Fornicata with Strikingly Simplified Ultrastructure. Journal of Eukaryotic Microbiology, 2017, 64, 422-433.	0.8	12
596	Spatio-temporal variability of periphytic protozoa related to environment in the Niyang River, Tibet, China. Chinese Journal of Oceanology and Limnology, 2017, 35, 489-500.	0.7	1
597	Medical Parasitology Taxonomy Update: January 2012 to December 2015. Journal of Clinical Microbiology, 2017, 55, 43-47.	1.8	15
598	An intact plastid genome is essential for the survival of colorless Euglena longa but not Euglena gracilis. Current Genetics, 2017, 63, 331-341.	0.8	17
599	A trophic framework for animal origins. Geobiology, 2017, 15, 197-210.	1.1	17
600	An evolutionary balance: conservation vs innovation in ciliate membrane trafficking. Traffic, 2017, 18, 18-28.	1.3	27
601	Metagenomics analysis reveals a new metallothionein family: Sequence and metal-binding features of new environmental cysteine-rich proteins. Journal of Inorganic Biochemistry, 2017, 167, 1-11.	1.5	35
602	Exploring the evolution of the proteins of the plant nuclear envelope. Nucleus, 2017, 8, 46-59.	0.6	46
603	<i>Didymium azorellae</i> , a new myxomycete from cushion plants of cold arid areas of South America. Mycologia, 2017, 109, 993-1002.	0.8	12
604	Cyanidiales: Evolution and Habitats. , 2017, , 3-15.		2

#	Article	IF	CITATIONS
605	Largeâ€scale patterns of biofilmâ€dwelling ciliate communities in a river network: Only small effects of stream order. International Review of Hydrobiology, 2017, 102, 114-124.	0.5	6
607	The Phylogeny of Myxomycetes. , 2017, , 83-106.		6
608	Taxonomy and Systematics: Current Knowledge and Approaches on the Taxonomic Treatment of Myxomycetes. , 2017, , 205-251.		13
609	Optics-based surveys of large unicellular zooplankton: a case study on radiolarians and phaeodarians. Plankton and Benthos Research, 2017, 12, 95-103.	0.2	15
610	First Reports on Heterotrophic Flagellates in the Mires of Arkhangelsk Region, Russia. Biology Bulletin, 2017, 44, 1007-1018.	0.1	2
611	Arctic Ocean. , 0, , 705-728.		0
612	What Defines the "Kingdom―Fungi?. , 2017, , 57-77.		6
613	Key Ecological Roles for Zoosporic True Fungi in Aquatic Habitats. , 2017, , 399-416.		1
614	Planktonic foraminifera-derived environmental DNA extracted from abyssal sediments preserves patterns of plankton macroecology. Biogeosciences, 2017, 14, 2741-2754.	1.3	36
615	The Role of Cytoplasmic mRNA Cap-Binding Protein Complexes in Trypanosoma brucei and Other Trypanosomatids. Pathogens, 2017, 6, 55.	1.2	52
616	Herbarium of the Pontifical Catholic University of Paraná (HUCP), Curitiba, Southern Brazil. Data, 2017, 2, 10.	1.2	1
617	Deep Transcriptome Sequencing of Two Green Algae, Chara vulgaris and Chlamydomonas reinhardtii, Provides No Evidence of Organellar RNA Editing. Genes, 2017, 8, 80.	1.0	24
618	Intercompartmental Piecewise Gene Transfer. Genes, 2017, 8, 260.	1.0	7
619	Microbial Eukaryotes in an Arctic Under-Ice Spring Bloom North of Svalbard. Frontiers in Microbiology, 2017, 8, 1099.	1.5	15
620	Retrotransposon Domestication and Control in Dictyostelium discoideum. Frontiers in Microbiology, 2017, 8, 1869.	1.5	6
621	Efficient Method for the Rapid Purification of <i>Nosema ceranae</i> Spores. Mycobiology, 2017, 45, 204-208.	0.6	2
622	The emerging picture of the mitochondrial protein import complexes of Amoebozoa supergroup. BMC Genomics, 2017, 18, 997.	1.2	7
623	InsP3 Signaling in Apicomplexan Parasites. Current Topics in Medicinal Chemistry, 2017, 17, 2158-2165.	1.0	49

#	Article	IF	CITATIONS
624	Dynamic species classification of microorganisms across time, abiotic and biotic environments—A sliding window approach. PLoS ONE, 2017, 12, e0176682.	1.1	21
625	Archigregarines of the English Channel revisited: New molecular data on Selenidium species including early described and new species and the uncertainties of phylogenetic relationships. PLoS ONE, 2017, 12, e0187430.	1.1	14
626	The ApaH-like phosphatase TbALPH1 is the major mRNA decapping enzyme of trypanosomes. PLoS Pathogens, 2017, 13, e1006456.	2.1	35
627	Motility in blastogregarines (Apicomplexa): Native and drug-induced organisation of Siedleckia nematoides cytoskeletal elements. PLoS ONE, 2017, 12, e0179709.	1.1	13
628	Rethinking the evolution of eukaryotic metabolism: novel cellular partitioning of enzymes in stramenopiles links serine biosynthesis to glycolysis in mitochondria. BMC Evolutionary Biology, 2017, 17, 241.	3.2	23
629	A synoptic overview of golden jackal parasites reveals high diversity of species. Parasites and Vectors, 2017, 10, 419.	1.0	41
630	Polyphasic insights into the microbiomes of the Takamatsuzuka Tumulus and Kitora Tumulus. Journal of General and Applied Microbiology, 2017, 63, 63-113.	0.4	22
631	Quest of Soil Protists in a New Era. Microbes and Environments, 2017, 32, 99-102.	0.7	6
632	Myosin repertoire expansion coincides with eukaryotic diversification in the Mesoproterozoic era. BMC Evolutionary Biology, 2017, 17, 211.	3.2	50
633	The Carboxy Terminus of YCF1 Contains a Motif Conserved throughout >500 Myr of Streptophyte Evolution. Genome Biology and Evolution, 2017, 9, 473-479.	1.1	14
634	Evolution of Life on Earth. , 2017, , 15-26.		0
635	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
636	Characterization of Cytosine Methylation and the DNA Methyltransferases of <i>Toxoplasma gondii</i> . International Journal of Biological Sciences, 2017, 13, 458-470.	2.6	13
637	The Evolutionary Landscape of Dbl-Like RhoCEF Families: Adapting Eukaryotic Cells to Environmental Signals. Genome Biology and Evolution, 2017, 9, 1471-1486.	1.1	47
638	Vertical distribution patterns of Radiolaria Polycystina (Protista) in the World Ocean: living ranges, isothermal submersion and settling shells. Journal of Plankton Research, 2017, 39, 330-349.	0.8	40
639	ESTRUTURA ESPACIAL E TEMPORAL DA COMUNIDADE MICROPLANCTA "NICA " "Tenenbaum, D.R., Nascimento, S.M, Vianna, S., Fragoso, G., Hatherly, M., Moraes, R. Estrutura espacial e temporal da comunidade microplanctĂ nica. 2017. In: FalcĂ£o, A.P.C., Moreira, D.L., editores. Ambiente pelĂ;gico: caracterizaĂ§Ă£o ambiental regional da Bacia de Campos, AtlĂ¢ntico Sudoeste. Rio de Janeiro: Elsevier. Habitats, v. 5. p.		1
640	Plastid Autonomy vs Nuclear Control Over Plastid Function. Advances in Botanical Research, 2018, 85, 1-28.	0.5	4
641	Microbial Diversity in the Eukaryotic SAR Clade: Illuminating the Darkness Between Morphology and Molecular Data. BioEssays, 2018, 40, e1700198.	1.2	43

#	Article	IF	CITATIONS
642	Diversity and evolution of four-domain voltage-gated cation channels of eukaryotes and their ancestral functional determinants. Scientific Reports, 2018, 8, 3539.	1.6	24
643	ChIP-ping the branches of the tree: functional genomics and the evolution of eukaryotic gene regulation. Briefings in Functional Genomics, 2018, 17, 116-137.	1.3	5
644	Benthic foraminifera in transitional environments in the English Channel and the southern North Sea: A proxy for regional-scale environmental and paleo-environmental characterisations. Marine Environmental Research, 2018, 137, 37-48.	1.1	25
645	Phylogeny and Morphology of New Diplonemids from Japan. Protist, 2018, 169, 158-179.	0.6	44
646	Multigene phylogeny and cell evolution of chromist infrakingdom Rhizaria: contrasting cell organisationAof sisterAphyla Cercozoa and Retaria. Protoplasma, 2018, 255, 1517-1574.	1.0	66
647	Extensive molecular tinkering in the evolution of the membrane attachment mode of the Rheb CTPase. Scientific Reports, 2018, 8, 5239.	1.6	9
648	Current view on phylogeny within the genus Flabellula Schaeffer, 1926 (Amoebozoa: Leptomyxida). European Journal of Protistology, 2018, 64, 40-53.	0.5	1
649	Discrepancies Between Molecular and Morphological Databases of Soil Ciliates Studied for Temperate Grasslands of Central Europe. Protist, 2018, 169, 521-538.	0.6	10
650	Introduction into Parasitic Protozoa. , 2018, , 1-10.		4
651	Babesia in Domestic Ruminants. , 2018, , 215-239.		6
652	Diversity and community composition of pico- and nanoplanktonic protists in the Vistula River estuary (Gulf of Gdańsk, Baltic Sea). Estuarine, Coastal and Shelf Science, 2018, 207, 242-249.	0.9	22
653	Molecular phylogeny and comparative morphology indicate that odontostomatids (Alveolata,) Tj ETQq1 1 0.7843 Evolution, 2018, 126, 382-389.	814 rgBT / 1.2	Overlock 10 25
654	Combined morphological and phylogenomic re-examination of malawimonads, a critical taxon for inferring the evolutionary history of eukaryotes. Royal Society Open Science, 2018, 5, 171707.	1.1	34
655	<scp>dinoref</scp> : A curated dinoflagellate (Dinophyceae) reference database for the 18S rRNA gene. Molecular Ecology Resources, 2018, 18, 974-987.	2.2	40
656	Regulation of early endosomes across eukaryotes: Evolution and functional homology of Vps9 proteins. Traffic, 2018, 19, 546-563.	1.3	12
657	Morphological Redescription of <i>Opalina undulata</i> Nie 1932 from <i>Fejervarya limnocharis</i> with Molecular Phylogenetic Study of Opalinids (Heterokonta, Opalinea). Journal of Eukaryotic Microbiology, 2018, 65, 783-791.	0.8	6
658	Fe–S cluster assembly in the supergroup Excavata. Journal of Biological Inorganic Chemistry, 2018, 23, 521-541.	1.1	17
659	Pervasive contingency and entrenchment in a billion years of Hsp90 evolution. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 4453-4458.	3.3	93

#	Article	IF	CITATIONS
660	Morphological, ontogenetic and molecular data support strongylidiids as being closely related to Dorsomarginalia (Protozoa, Ciliophora) and reactivation of the family Strongylidiidae Fauré-Fremiet, 1961. Zoological Journal of the Linnean Society, 2018, 184, 237-254.	1.0	38
661	Aberrant and accidental trichomonad flagellate infections: rare or underdiagnosed?. Transactions of the Royal Society of Tropical Medicine and Hygiene, 2018, 112, 64-72.	0.7	9
662	Reconstruction of the sialylation pathway in the ancestor of eukaryotes. Scientific Reports, 2018, 8, 2946.	1.6	20
663	Parasitische Protozoen. , 2018, , 99-247.		0
664	Plastid phylogenomics with broad taxon sampling further elucidates the distinct evolutionary origins and timing of secondary green plastids. Scientific Reports, 2018, 8, 1523.	1.6	66
665	Opportunistic but Lethal: The Mystery of Paramoebae. Trends in Parasitology, 2018, 34, 404-419.	1.5	41
666	Soil protists: a fertile frontier in soil biology research. FEMS Microbiology Reviews, 2018, 42, 293-323.	3.9	368
667	Classification of Fungi. , 2018, , 1229-1231.e1.		1
668	Distribution and Evolution of Peroxisomes in Alveolates (Apicomplexa, Dinoflagellates, Ciliates). Genome Biology and Evolution, 2018, 10, 1-13.	1.1	21
669	<i>Sappinia</i> sp. (Amoebozoa: Thecamoebida) and <i>Rosculus</i> sp. (SAR: Cercozoa) Isolated From King Penguin Guano Collected in the Subantarctic (South Georgia, Salisbury Plain) and their Coexistence in Culture. Journal of Eukaryotic Microbiology, 2018, 65, 544-555.	0.8	4
670	Molecular Phylogenetic Positions and Ultrastructure of Marine Gregarines (Apicomplexa) <i>Cuspisella ishikariensis</i> n. gen., n. sp. and <i>Loxomorpha</i> cf. <i>harmothoe</i> from Western Pacific scaleworms (Polynoidae). Journal of Eukaryotic Microbiology, 2018, 65, 637-647.	0.8	3
671	TelAP1 links telomere complexes with developmental expression site silencing in African trypanosomes. Nucleic Acids Research, 2018, 46, 2820-2833.	6.5	24
672	Intrageneric Variability Between the Chloroplast Genomes of <i>Trachelomonas grandis</i> and <i>Trachelomonas volvocina</i> and Phylogenomic Analysis of Phototrophic Euglenoids. Journal of Eukaryotic Microbiology, 2018, 65, 648-660.	0.8	3
673	An Orphan Protist <i>Quadricilia rotundata</i> Finally Finds Its Phylogenetic Home in Cercozoa. Journal of Eukaryotic Microbiology, 2018, 65, 729-732.	0.8	3
674	Labyrinthulomycota from Brazilian mangrove swamps and coastal waters. Botanica Marina, 2018, 61, 65-74.	0.6	3
675	Phylogenomics Places Orphan Protistan Lineages in a Novel Eukaryotic Super-Group. Genome Biology and Evolution, 2018, 10, 427-433.	1.1	112
676	Further insights into the highly derived haptorids (Ciliophora, Litostomatea): Phylogeny based on multigene data. Zoologica Scripta, 2018, 47, 231-242.	0.7	33
677	The F ₁ â€ <scp>ATP</scp> ase from <i>Trypanosoma brucei</i> is elaborated by three copies of an additional p18â€subunit. FEBS Journal, 2018, 285, 614-628.	2.2	20

#	Article	IF	CITATIONS
678	A single-cysteine mutant and chimeras of essential Leishmania Erv can complement the loss of Erv1 but not of Mia40 in yeast. Redox Biology, 2018, 15, 363-374.	3.9	12
679	Neobodonids are dominant kinetoplastids in the global ocean. Environmental Microbiology, 2018, 20, 878-889.	1.8	27
680	200 years of marine research at Senckenberg: selected highlights. Marine Biodiversity, 2018, 48, 159-178.	0.3	4
681	Plastid Genomes in the Myzozoa. Advances in Botanical Research, 2018, 85, 55-94.	0.5	4
682	Identification of a novel fused gene family implicates convergent evolution in eukaryotic calcium signaling. BMC Genomics, 2018, 19, 306.	1.2	4
683	Effect of ammonium and high light intensity on the accumulation of lipids in Nannochloropsis oceanica (CCAP 849/10) and Phaeodactylum tricornutum (CCAP 1055/1). Biotechnology for Biofuels, 2018, 11, 60.	6.2	28
684	Transformation of <i>Diplonema papillatum</i> , the type species of the highly diverse and abundant marine microeukaryotes Diplonemida (Euglenozoa). Environmental Microbiology, 2018, 20, 1030-1040.	1.8	20
685	First Ultrastructural and Molecular Phylogenetic Evidence from the Blastogregarines, an Early Branching Lineage of Plesiomorphic Apicomplexa. Protist, 2018, 169, 697-726.	0.6	14
686	Plastid Transcript Editing across Dinoflagellate Lineages Shows Lineage-Specific Application but Conserved Trends. Genome Biology and Evolution, 2018, 10, 1019-1038.	1.1	22
687	Functional Analysis of Hif1 Histone Chaperone in <i>Saccharomyces cerevisiae</i> . G3: Genes, Genomes, Genetics, 2018, 8, 1993-2006.	0.8	8
688	Identification and characterisation of the cryptic Golgi apparatus in <i>Naegleria gruberi</i> . Journal of Cell Science, 2018, 131, .	1.2	6
689	Integrative analysis of large scale transcriptome data draws a comprehensive landscape of Phaeodactylum tricornutum genome and evolutionary origin of diatoms. Scientific Reports, 2018, 8, 4834.	1.6	131
690	Short-term dynamics and interactions of marine protist communities during the spring–summer transition. ISME Journal, 2018, 12, 1907-1917.	4.4	84
691	Transepithelial migration of mucosal hemocytes in Crassostrea virginica and potential role in Perkinsus marinus pathogenesis. Journal of Invertebrate Pathology, 2018, 153, 122-129.	1.5	18
692	Permian smaller foraminifers: taxonomy, biostratigraphy and biogeography. Geological Society Special Publication, 2018, 450, 205-252.	0.8	15
693	Cytochrome P450 diversity in the tree of life. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2018, 1866, 141-154.	1.1	229
694	Microeukaryote community in a partial nitritation reactor prior to anammox and an insight into the potential of ciliates as performance bioindicators. New Biotechnology, 2018, 43, 3-12.	2.4	5
695	Morphology and Phylogeny of Two Novel Ciliates, <i>Arcanisutura chongmingensis</i> n. gen., n. sp. and <i>Naxella paralucida</i> n. sp. from Shanghai, China. Journal of Eukaryotic Microbiology, 2018, 65, 48-60.	0.8	8

#	Article	IF	CITATIONS
696	Methodological advances to study the diversity of soil protists and their functioning in soil food webs. Applied Soil Ecology, 2018, 123, 328-333.	2.1	62
697	Tonian (Neoproterozoic) eukaryotic and prokaryotic organic-walled microfossils from the upper VisingsA¶ Group, Sweden. Palynology, 2018, 42, 220-254.	0.7	30
698	Not in your usual Top 10: protists that infect plants and algae. Molecular Plant Pathology, 2018, 19, 1029-1044.	2.0	55
699	Reductive evolution of chloroplasts in non-photosynthetic plants, algae and protists. Current Genetics, 2018, 64, 365-387.	0.8	81
700	Potential contribution of surface-dwelling Sargassum algae to deep-sea ecosystems in the southern North Atlantic. Deep-Sea Research Part II: Topical Studies in Oceanography, 2018, 148, 21-34.	0.6	37
701	Variation in the SSUrDNA of the Genus <i>Protostelium</i> Leads to a New Phylogenetic Understanding of the Genus and of the Species Concept for <i>Protostelium mycophaga</i> (Protosteliida, Amoebozoa). Journal of Eukaryotic Microbiology, 2018, 65, 331-344.	0.8	9
702	Seeing the endomembrane system for the trees: Evolutionary analysis highlights the importance of plants as models for eukaryotic membrane-trafficking. Seminars in Cell and Developmental Biology, 2018, 80, 142-152.	2.3	17
703	A proteomic portrait of dinoflagellate chromatin reveals abundant RNA-binding proteins. Chromosoma, 2018, 127, 29-43.	1.0	13
704	Evidence of the supercomplex organization of photosystem II and light-harvesting complexes in Nannochloropsis granulata. Photosynthesis Research, 2018, 136, 49-61.	1.6	13
705	The diversity and biogeography of abundant and rare intertidal marine microeukaryotes explained by environment and dispersal limitation. Environmental Microbiology, 2018, 20, 462-476.	1.8	112
706	Molecular Phylogenetic Positions of Two New Marine Gregarines (Apicomplexa)— <i>Paralecudina anankea</i> n. sp. and <i>Lecudina caspera</i> n. sp.—from the Intestine of <i>Lumbrineris inflata</i> (Polychaeta) Show Patterns of Coâ€evolution. Journal of Eukaryotic Microbiology, 2018, 65, 211-219.	0.8	4
707	Vase-shaped microfossil biostratigraphy with new data from Tasmania, Svalbard, Greenland, Sweden and the Yukon. Precambrian Research, 2018, 319, 19-36.	1.2	42
708	Barcoding in trypanosomes. Parasitology, 2018, 145, 563-573.	0.7	11
709	Phylogenetic characterization of transporter proteins in the cnidarian-dinoflagellate symbiosis. Molecular Phylogenetics and Evolution, 2018, 120, 307-320.	1.2	30
710	Novel Diversity of Deeply Branching Holomycota and Unicellular Holozoans Revealed by Metabarcoding in Middle ParanÃ _i River, Argentina. Frontiers in Ecology and Evolution, 2018, 6, .	1.1	20
711	On the Systematics and Scale for Measuring Historical Pathways of the Development of Morphological Diversity. Paleontological Journal, 2018, 52, 1789-1798.	0.2	0
712	Phylogenetic taxon definitions for Fungi, Dikarya, Ascomycota and Basidiomycota. IMA Fungus, 2018, 9, 291-298.	1.7	26
714	NommPred: Prediction of Mitochondrial and Mitochondrion-Related Organelle Proteins of Nonmodel Organisms. Evolutionary Bioinformatics, 2018, 14, 117693431881983.	0.6	17

#	Article	IF	CITATIONS
715	Comparative Genomics Supports Sex and Meiosis in Diverse Amoebozoa. Genome Biology and Evolution, 2018, 10, 3118-3128.	1.1	25
716	Phylogeny and Classification of Novel Diversity in Sainouroidea (Cercozoa, Rhizaria) Sheds Light on a Highly Diverse and Divergent Clade. Protist, 2018, 169, 853-874.	0.6	5
717	Advanced Techniques in Diagnostic Parasitology. , 2018, , 199-218.		1
718	A single class of ARF GTPase activated by several pathway-specific ARF-GEFs regulates essential membrane traffic in Arabidopsis. PLoS Genetics, 2018, 14, e1007795.	1.5	28
719	Order of removal of conventional and nonconventional introns from nuclear transcripts of Euglena gracilis. PLoS Genetics, 2018, 14, e1007761.	1.5	14
720	Common ancestry of heterodimerizing TALE homeobox transcription factors across Metazoa and Archaeplastida. BMC Biology, 2018, 16, 136.	1.7	21
721	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
722	Comparative Biology of Centrosomal Structures in Eukaryotes. Cells, 2018, 7, 202.	1.8	6
723	Nuclear genome sequence of the plastid-lacking cryptomonad Goniomonas avonlea provides insights into the evolution of secondary plastids. BMC Biology, 2018, 16, 137.	1.7	42
724	Perfection of eccentricity: Mitochondrial genomes of diplonemids. IUBMB Life, 2018, 70, 1197-1206.	1.5	24
725	Branched late-steps of the cytosolic iron-sulphur cluster assembly machinery of Trypanosoma brucei. PLoS Pathogens, 2018, 14, e1007326.	2.1	2
726	Heavy Metal Pumps in Plants: Structure, Function and Origin. Advances in Botanical Research, 2018, , 57-89.	0.5	11
727	Epiplasts: Membrane Skeletons and Epiplastin Proteins in Euglenids, Glaucophytes, Cryptophytes, Ciliates, Dinoflagellates, and Apicomplexans. MBio, 2018, 9, .	1.8	23
728	Spatial Variability of Picoeukaryotic Communities in the Mariana Trench. Scientific Reports, 2018, 8, 15357.	1.6	31
729	Glutamate synthases from conifers: gene structure and phylogenetic studies. BMC Genomics, 2018, 19, 65.	1.2	11
730	Protist species richness and soil microbiome complexity increase towards climax vegetation in the Brazilian Cerrado. Communications Biology, 2018, 1, 135.	2.0	58
731	A putative NEM1 homologue regulates lipid droplet biogenesis via PAH1 in Tetrahymena thermophila. Journal of Biosciences, 2018, 43, 693-706.	0.5	6
732	Fe–S Cluster Assembly in Oxymonads and Related Protists. Molecular Biology and Evolution, 2018, 35, 2712-2718.	3.5	19

#	Article	IF	CITATIONS
733	Dynamics and determinants of amoeba community, occurrence and abundance in subtropical reservoirs and rivers. Water Research, 2018, 146, 177-186.	5.3	44
734	Transcriptional responses of the marine green alga <i>Micromonas pusilla</i> and an infecting prasinovirus under different phosphate conditions. Environmental Microbiology, 2018, 20, 2898-2912.	1.8	25
735	Ancestral State Reconstruction of the Apoptosis Machinery in the Common Ancestor of Eukaryotes. G3: Genes, Genomes, Genetics, 2018, 8, 2121-2134.	0.8	32
736	<i>Burkholderia</i> bacteria use chemotaxis to find social amoeba <i>Dictyostelium discoideum</i> hosts. ISME Journal, 2018, 12, 1977-1993.	4.4	41
737	Bodenorganismen und ihr Lebensraum. , 2018, , 103-149.		0
738	What Makes an Animal? The Molecular Quest for the Origin of the Animal Kingdom. Integrative and Comparative Biology, 2018, 58, 654-665.	0.9	15
739	High-level classification of the Fungi and a tool for evolutionary ecological analyses. Fungal Diversity, 2018, 90, 135-159.	4.7	450
740	Is Myxomycetes (Amoebozoa) a Truly Ambiregnal Group? A Major Issue in Protist Nomenclature. Protist, 2018, 169, 484-493.	0.6	4
741	Decoupling of Nuclear Division Cycles and Cell Size during the Coenocytic Growth of the Ichthyosporean Sphaeroforma arctica. Current Biology, 2018, 28, 1964-1969.e2.	1.8	27
742	Global characterization of the Dicer-like protein DrnB roles in miRNA biogenesis in the social amoeba <i>Dictyostelium discoideum</i> . RNA Biology, 2018, 15, 937-954.	1.5	9
743	Lost in the Light: Plastid Genome Evolution in Nonphotosynthetic Algae. Advances in Botanical Research, 2018, 85, 29-53.	0.5	2
744	When Less is More: Red Algae as Models for Studying Gene Loss and Genome Evolution in Eukaryotes. Critical Reviews in Plant Sciences, 2018, 37, 81-99.	2.7	30
745	Evaluating rodent experimental models for studies of Blastocystis ST1. Experimental Parasitology, 2018, 191, 55-61.	0.5	11
746	Comparative Plastid Genomics of Glaucophytes. Advances in Botanical Research, 2018, 85, 95-127.	0.5	6
747	Pythium insidiosum isolated from infected mosquito larvae in central Brazil. Acta Tropica, 2018, 185, 344-348.	0.9	13
748	Phylogenetic relationship analyses of complicated class Spirotrichea based on transcriptomes from three diverse microbial eukaryotes: Uroleptopsis citrina, Euplotes vannus and Protocruzia tuzeti. Molecular Phylogenetics and Evolution, 2018, 129, 338-345.	1.2	36
749	Mitochondrial Glycolysis in a Major Lineage of Eukaryotes. Genome Biology and Evolution, 2018, 10, 2310-2325.	1.1	62
750	Centrosome Remodelling in Evolution. Cells, 2018, 7, 71.	1.8	46

ARTICLE IF CITATIONS # Apicomplexa Cell Cycles: Something Old, Borrowed, Lost, and New. Trends in Parasitology, 2018, 34, 751 1.5 54 759-771. Diversity and Evolutionary History of the Symbiontida (Euglenozoa). Frontiers in Ecology and 1.1 Evolution, 2018, 6, . Trypanosomal mitochondrial intermediate peptidase does not behave as a classical mitochondrial 753 9 1.1 processing peptidase. PLoS ONE, 2018, 13, e0196474. Cell polarity: having and making sense of directionâ€" on the evolutionary significance of the primary 754 cilium/centrosome organ in Metazoa. Open Biology, 2018, 8, . Competition between Silicifiers and Non-silicifiers in the Past and Present Ocean and Its Evolutionary 755 1.2 29 Impacts. Frontiers in Marine Science, 2018, 5, . Eukaryotes Pose New Problems., 2018, , 117-133. Evolution of protein trafficking in kinetoplastid parasites: Complexity and pathogenesis. Traffic, 2018, 757 1.3 8 19,803-812. Microalgal Systematics., 2018,, 73-107. 758 Ticks and Tick-Borne Infections: Complex Ecology, Agents, and Host Interactions. Veterinary Sciences, 759 0.6 105 2018, 5, 60. Vermamoeba vermiformis: a Free-Living Amoeba of Interest. Microbial Ecology, 2018, 76, 991-1001. 1.4 EukRefâ€Ciliophora: a manually curated, phylogenyâ€based database of small subunit rRNA gene sequences 761 1.8 27 of ciliates. Environmental Microbiology, 2018, 20, 2218-2230. Keap1–MCM3 interaction is a potential coordinator of molecular machineries of antioxidant response 24 and genomic DNA replication in metazoa. Scientific Reports, 2018, 8, 12136. Microbiome of a revegetated iron-mining site and pristine ecosystems from the Brazilian Cerrado. 764 2.1 21 Applied Soil Ecology, 2018, 131, 55-65. Molecular survey of coccidian infections of the side-blotched lizard Uta stansburiana on San Benito 0.8 Oeste Island, México. Parasite, 2018, 25, 43. Nutritional Intake by Ectoplasmic Nets of Schizochytrium aggregatum (Labyrinthulomycetes,) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 182 766 Protozoa â[~]†., 2018,,. Evolutionary cell biology traces the rise of the exomer complex in Fungi from an ancient eukaryotic 768 1.6 7 component. Scientific Reports, 2018, 8, 11154.

CITATION REPORT

769Animal origins and the Tonian Earth system. Emerging Topics in Life Sciences, 2018, 2, 289-298.1.112

#	Article	IF	CITATIONS
770	Molecular and Morphological Diversity of the Oxymonad Genera Monocercomonoides and Blattamonas gen. nov Protist, 2018, 169, 744-783.	0.6	19
771	Foraminiferal growth and test development. Earth-Science Reviews, 2018, 185, 140-162.	4.0	15
772	Diatom Allantoin Synthase Provides Structural Insights into Natural Fusion Protein Therapeutics. ACS Chemical Biology, 2018, 13, 2237-2246.	1.6	5
773	Benthic ciliate diversity and community composition along water depth gradients: a comparison between the intertidal and offshore areas. European Journal of Protistology, 2018, 65, 31-41.	0.5	5
774	Life Cycle, Ultrastructure, and Phylogeny of New Diplonemids and Their Endosymbiotic Bacteria. MBio, 2018, 9, .	1.8	50
775	Definition of benthic foraminiferal bioprovinces in transitional environments of the Eastern English Channel and the Southern North Sea. Revue De Micropaleontologie, 2018, 61, 223-234.	0.8	10
776	Symbiotic flagellate protists as cryptic drivers of adaptation and invasiveness of the subterranean termite <i>Reticulitermes grassei</i> Clément. Ecology and Evolution, 2018, 8, 5242-5253.	0.8	10
777	Mitochondrial RNA Editing and Processing in Diplonemid Protists. Nucleic Acids and Molecular Biology, 2018, , 145-176.	0.2	5
778	Distinct patterns and processes of abundant and rare eukaryotic plankton communities following a reservoir cyanobacterial bloom. ISME Journal, 2018, 12, 2263-2277.	4.4	412
779	Recent advances in trypanosomatid research: genome organization, expression, metabolism, taxonomy and evolution. Parasitology, 2019, 146, 1-27.	0.7	121
780	Description of Armaparvus languidus n. gen. n. sp. Confirms Ultrastructural Unity of Cutosea (Amoebozoa, Evosea). Journal of Eukaryotic Microbiology, 2019, 66, 158-166.	0.8	4
781	<i>Dactylomonas</i> gen. nov., a Novel Lineage of Heterolobosean Flagellates with Unique Ultrastructure, Closely Related to the Amoeba <i>Selenaion koniopes</i> Park, De Jonckheere & Simpson, 2012. Journal of Eukaryotic Microbiology, 2019, 66, 120-139.	0.8	12
782	There Is Treasure Everywhere: Reductive Plastid Evolution in Apicomplexa in Light of Their Close Relatives. Biomolecules, 2019, 9, 378.	1.8	29
783	Phytoplankton growth characterization in short term MPN culture assays using 18S metabarcoding and qRT-PCR. Water Research, 2019, 164, 114941.	5.3	4
784	Evolution of the Cholesterol Biosynthesis Pathway in Animals. Molecular Biology and Evolution, 2019, 36, 2548-2556.	3.5	37
785	Evidence of conditioned behavior in amoebae. Nature Communications, 2019, 10, 3690.	5.8	30
786	Alternative splicing is required for stage differentiation in malaria parasites. Genome Biology, 2019, 20, 151.	3.8	29
788	Further analyses on the phylogeny of the subclass Scuticociliatia (Protozoa, Ciliophora) based on both nuclear and mitochondrial data. Molecular Phylogenetics and Evolution, 2019, 139, 106565.	1.2	33

#	Article	IF	Citations
790	Phytophthora infestans Dihydroorotate Dehydrogenase Is a Potential Target for Chemical Control – A Comparison With the Enzyme From Solanum tuberosum. Frontiers in Microbiology, 2019, 10, 1479.	1.5	12
791	The Med31 Conserved Component of the Divergent Mediator Complex in Tetrahymena thermophila Participates in Developmental Regulation. Current Biology, 2019, 29, 2371-2379.e6.	1.8	13
792	Nitroreductases of bacterial origin in Giardia lamblia : Potential role in detoxification of xenobiotics. MicrobiologyOpen, 2019, 8, e904.	1.2	8
793	Comparative Pathobiology of the Intestinal Protozoan Parasites Giardia lamblia, Entamoeba histolytica, and Cryptosporidium parvum. Pathogens, 2019, 8, 116.	1.2	46
794	Introduction to Genome Biology and Diversity. Methods in Molecular Biology, 2019, 1910, 3-31.	0.4	8
796	Changes of Gene Expression in Euglena gracilis Obtained During the 29th DLR Parabolic Flight Campaign. Scientific Reports, 2019, 9, 14260.	1.6	10
797	Fungal evolution: diversity, taxonomy and phylogeny of the Fungi. Biological Reviews, 2019, 94, 2101-2137.	4.7	191
798	Successional trophic complexity and biogeographical structure of eukaryotic communities in waterworks' rapid sand filters. FEMS Microbiology Ecology, 2019, 95, .	1.3	3
799	Haemocystidium spp., a species complex infecting ancient aquatic turtles of the family Podocnemididae: First report of these parasites in Podocnemis vogli from the Orinoquia. International Journal for Parasitology: Parasites and Wildlife, 2019, 10, 299-309.	0.6	7
800	Thecamoeba cosmophorea n. sp. (Amoebozoa, Discosea, Thecamoebida) — An example of sibling species within the genus Thecamoeba. European Journal of Protistology, 2019, 67, 132-141.	0.5	7
801	The Biochemistry and Evolution of the Dinoflagellate Nucleus. Microorganisms, 2019, 7, 245.	1.6	29
802	Assortative Mating in Animals and Its Role for Speciation. American Naturalist, 2019, 194, 865-875.	1.0	35
803	Heterotrophic Flagellates from Sphagnum Bogs and Terrace-Forest and Floodplain Water Bodies of the Central Russian Forest-Steppe. Inland Water Biology, 2019, 12, 276-289.	0.2	0
804	Analysis of Microbial Community Dynamics during the Acclimatization Period of a Membrane Bioreactor Treating Table Olive Processing Wastewater. Applied Sciences (Switzerland), 2019, 9, 3647.	1.3	4
805	Meiofaunal diversity in the Atlantic Forest soil: A quest for nematodes in a native reserve using eukaryotic metabarcoding analysis. Forest Ecology and Management, 2019, 453, 117591.	1.4	18
806	Microscopical Studies on Ministeria vibrans Tong, 1997 (Filasterea) Highlight the Cytoskeletal Structure of the Common Ancestor of Filasterea, Metazoa and Choanoflagellata. Protist, 2019, 170, 385-396.	0.6	15
807	Redescription of Opalina triangulata (Heterokonta, Opalinea) from Fejervarya limnocharis based on morphological and molecular data. European Journal of Protistology, 2019, 71, 125639.	0.5	3
808	Gene identification and functional characterization of a Δ12 fatty acid desaturase in Tetrahymena thermophila and its influence in homeoviscous adaptation to low temperature. Biochimica Et Biophysica Acta - Molecular and Cell Biology of Lipids, 2019, 1864, 1644-1655.	1.2	6

	CITATIO	CITATION REPORT	
#	Article	IF	Citations
809	Using single-cell transcriptomics to understand functional states and interactions in microbial eukaryotes. Philosophical Transactions of the Royal Society B: Biological Sciences, 2019, 374, 20190098.	1.8	20
810	Nutritional intake of Aplanochytrium (Labyrinthulea, Stramenopiles) from living diatoms revealed by culture experiments suggesting the new prey–predator interactions in the grazing food web of the marine ecosystem. PLoS ONE, 2019, 14, e0208941.	1.1	28
811	Plastid Genomes from Diverse Glaucophyte Genera Reveal a Largely Conserved Gene Content and Limited Architectural Diversity. Genome Biology and Evolution, 2019, 11, 174-188.	1.1	16
812	Calcium negatively regulates secretion from dense granules in <scp> <i>Toxoplasma gondii</i> </scp> . Cellular Microbiology, 2019, 21, e13011.	1.1	18
813	Isolation and characterization of a tandem-repeated cysteine protease from the symbiotic dinoflagellate Symbiodinium sp. KB8. PLoS ONE, 2019, 14, e0211534.	1.1	2
814	Advances in isolation and preservation strategies of ecologically important marine protists, the thraustochytrids. , 2019, , 485-500.		2
815	Algae. , 2019, , 95-120.		2
816	Eukaryotes. , 2019, , 155-231.		0
817	Regulation of gene expression in trypanosomatids: living with polycistronic transcription. Open Biology, 2019, 9, 190072.	1.5	166
818	Microalgae for High-Value Products Towards Human Health and Nutrition. Marine Drugs, 2019, 17, 304.	2.2	355
819	Horizontal and endosymbiotic gene transfer in early plastid evolution. New Phytologist, 2019, 224, 618-624.	3.5	57
820	Plasmodium pseudo-Tyrosine Kinase-like binds PP1 and SERA5 and is exported to host erythrocytes. Scientific Reports, 2019, 9, 8120.	1.6	9
821	The Chloroplast and Photosynthetic Eukaryotes. , 2019, , 269-310.		2
822	A eukaryotic community succession based method for postmortem interval (PMI) estimation of decomposing porcine remains. Forensic Science International, 2019, 302, 109838.	1.3	17
823	The Cryptosporidium parvum gp60 glycoprotein expressed in the ciliate Tetrahymena thermophila is immunoreactive with sera of calves infected with Cryptosporidium oocysts. Veterinary Parasitology, 2019, 271, 45-50.	0.7	6
824	Analysis of an improved Cyanophora paradoxa genome assembly. DNA Research, 2019, 26, 287-299.	1.5	35
825	Incubation and grazing effects on spirotrich ciliate diversity inferred from molecular analyses of microcosm experiments. PLoS ONE, 2019, 14, e0215872.	1.1	9
826	Protozoan predation of <i>Escherichia coli</i> in hydroponic media of leafy vegetables. Soil Science and Plant Nutrition, 2019, 65, 234-242.	0.8	1

#	Article	IF	CITATIONS
827	Development of an autonomous biosampler to capture in situ aquatic microbiomes. PLoS ONE, 2019, 14, e0216882.	1.1	13
828	An online resource for marine fungi. Fungal Diversity, 2019, 96, 347-433.	4.7	133
829	Ediacaran microfossils from the Doushantuo Formation chert nodules in the Yangtze Gorges area, South China, and new biozones. Fossils and Strata, 2019, , 1-172.	2.0	31
831	Different Degrees of Niche Differentiation for Bacteria, Fungi, and Myxomycetes Within an Elevational Transect in the German Alps. Microbial Ecology, 2019, 78, 764-780.	1.4	16
832	A dynamically interacting flexible loop assists oligomerisation of the Caenorhabditis elegans centriolar protein SAS-6. Scientific Reports, 2019, 9, 3526.	1.6	3
833	Ultrastructure and phylogenetic characterization of the microsporidian parasite Heterosporis lessepsianus n. sp. (Microsporidia: Glugeidae) infecting the lizardfish Saurida lessepsianus (Pisces:) Tj ETQq1 1 C	.78 143 14 r	gBħ/Overlock
834	Evolutionary dynamics of the chromatophore genome in three photosynthetic Paulinella species. Scientific Reports, 2019, 9, 2560.	1.6	30
835	Comparative genomics reveals the unique evolutionary status of Plasmodiophora brassicae and the essential role of GPCR signaling pathways. Phytopathology Research, 2019, 1, .	0.9	17
836	Cell biology in phytopathogenic fungi during host infection: commonalities and differences. Journal of General Plant Pathology, 2019, 85, 163-173.	0.6	13
837	<i>Chrysochromulina andersonii sp. nov</i> . (Prymnesiophyceae), a new flagellate haptophyte symbiotic with radiolarians. Phycologia, 2019, 58, 211-224.	0.6	2
838	Proteomic Analysis of Histones H2A/H2B and Variant Hv1 in Tetrahymena thermophila Reveals an Ancient Network of Chaperones. Molecular Biology and Evolution, 2019, 36, 1037-1055.	3.5	12
839	Evolution and a revised nomenclature of P4 ATPases, a eukaryotic family of lipid flippases. Biochimica Et Biophysica Acta - Biomembranes, 2019, 1861, 1135-1151.	1.4	46
840	Transcriptome, proteome and draft genome of Euglena gracilis. BMC Biology, 2019, 17, 11.	1.7	98
841	Nuclear localization signals in trypanosomal proteins. Molecular and Biochemical Parasitology, 2019, 229, 15-23.	0.5	8
842	Changes in Trophic Groups of Protists With Conversion of Rainforest Into Rubber and Oil Palm Plantations. Frontiers in Microbiology, 2019, 10, 240.	1.5	48
843	Contrasted Micro-Eukaryotic Diversity Associated with Sphagnum Mosses in Tropical, Subtropical and Temperate Climatic Zones. Microbial Ecology, 2019, 78, 714-724.	1.4	11
844	Cells and Organisms. , 2019, , 2-37.		0
845	Morphological stasis in the first myxomycete from the Mesozoic, and the likely role of cryptobiosis. Scientific Reports, 2019, 9, 19730.	1.6	8

#	Article	IF	CITATIONS
846	tRNA functional signatures classify plastids as late-branching cyanobacteria. BMC Evolutionary Biology, 2019, 19, 224.	3.2	5
847	Responses of unicellular predators to cope with the phototoxicity of photosynthetic prey. Nature Communications, 2019, 10, 5606.	5.8	11
848	Nutrient sensing-the key to fungal p53-like transcription factors?. Fungal Genetics and Biology, 2019, 124, 8-16.	0.9	9
849	Dictyostelium: An Important Source of Structural and Functional Diversity in Drug Discovery. Cells, 2019, 8, 6.	1.8	24
850	Phylogenomics of Thecamoebida (Discosea, Amoebozoa) with the Description of Stratorugosa tubuloviscum gen. nov. sp. nov., a Freshwater Amoeba with a Perinuclear MTOC. Protist, 2019, 170, 8-20.	0.6	9
851	Biotechnology in ciliates: an overview. Critical Reviews in Biotechnology, 2019, 39, 220-234.	5.1	7
852	Protists: Puppet Masters of the Rhizosphere Microbiome. Trends in Plant Science, 2019, 24, 165-176.	4.3	215
853	Was the Mitochondrion Necessary to Start Eukaryogenesis?. Trends in Microbiology, 2019, 27, 96-104.	3.5	42
854	Community of dark-spored myxomycetes in ground litter and soil of taiga forest (Nizhne-Svirskiy) Tj ETQq0 0 0 rg	gBT /Overla	ock 10 Tf 50
855	Three monophyletic clusters in Retortamonas species isolated from vertebrates. Parasitology International, 2019, 69, 93-98.	0.6	2
856	The complex enzymology of mRNA decapping: Enzymes of four classes cleave pyrophosphate bonds. Wiley Interdisciplinary Reviews RNA, 2019, 10, e1511.	3.2	31
857	Swarming and Aggregation in the Parasitic Diplomonad Flagellate Spironucleus vortens. Journal of Eukaryotic Microbiology, 2019, 66, 545-552.	0.8	0
858	Epigenetics in the early divergent eukaryotic Giardia duodenalis: An update. Biochimie, 2019, 156, 123-128.	1.3	18
859	A comprehensive review on past, present and future aspects of canine theileriosis. Microbial Pathogenesis, 2019, 126, 116-122.	1.3	8
860	Does specialization imply rare fossil records of some benthic foraminifera: Late Palaeocene examples from the eastern Neo-Tethys (Meghalaya, NE India). Palaeogeography, Palaeoclimatology, Palaeoecology, 2019, 514, 124-134.	1.0	8
861	Revisions to the Classification, Nomenclature, and Diversity of Eukaryotes. Journal of Eukaryotic Microbiology, 2019, 66, 4-119.	0.8	904
862	Palaeobiology of latest Ediacaran phosphorites from the upper Khesen Formation, Khuvsgul Group, northern Mongolia. Journal of Systematic Palaeontology, 2019, 17, 501-532.	0.6	24

863	Pellicle ultrastructure demonstrates that <i>Moyeria</i> is a fossil euglenid. Palynology, 2020, 44, 461-471.	0.7	10
-----	---	-----	----

#	Article	IF	CITATIONS
864	The New Tree of Eukaryotes. Trends in Ecology and Evolution, 2020, 35, 43-55.	4.2	537
865	Life Cycles of Myxogastria Stemonitopsis typhina and Stemonitis fusca on Agar Culture. Journal of Eukaryotic Microbiology, 2020, 67, 66-75.	0.8	4

866
866
866
866
867
868
868
868
868
868
869
869
869
869
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860
860

867	Gene expression analysis of <i>Cyanophora paradoxa</i> reveals conserved abiotic stress responses between basal algae and flowering plants. New Phytologist, 2020, 225, 1562-1577.	3.5	10
868	Evidence of nuclear transport mechanisms in the protozoan parasite Giardia lamblia. Biochimica Et Biophysica Acta - Molecular Cell Research, 2020, 1867, 118566.	1.9	4
869	Evaluation of accuracy and precision in an amplicon sequencing workflow for marine protist communities. Limnology and Oceanography: Methods, 2020, 18, 20-40.	1.0	18
870	Genomeâ€wide Transcriptional Analysis ofTetrahymena thermophilaResponse to Exogenous Cholesterol. Journal of Eukaryotic Microbiology, 2020, 67, 209-222.	0.8	4
871	Is pallial mucus involved in Ostrea edulis defenses against the parasite Bonamia ostreae?. Journal of Invertebrate Pathology, 2020, 169, 107259.	1.5	10
872	Emergence and Evolution of ERM Proteins and Merlin in Metazoans. Genome Biology and Evolution, 2020, 12, 3710-3724.	1.1	9
873	The planktonic protist interactome: where do we stand after a century of research?. ISME Journal, 2020, 14, 544-559.	4.4	111
874	The Lipid Composition of Euglena gracilis Middle Plastid Membrane Resembles That of Primary Plastid Envelopes. Plant Physiology, 2020, 184, 2052-2063.	2.3	3
875	Composition and Function of Telomerase—A Polymerase Associated with the Origin of Eukaryotes. Biomolecules, 2020, 10, 1425.	1.8	16
876	The microwave assisted extraction sway on the features of antioxidant compounds and gelling biopolymers from Mastocarpus stellatus. Algal Research, 2020, 51, 102081.	2.4	37
877	Anti-nosemosis activity of phenolic compounds derived from <i>Artemisia dubia</i> and <i>Aster scaber</i> . Journal of Apicultural Research, 2022, 61, 519-529.	0.7	6
878	Adaption of microbial communities to the hostile environment in the Doce River after the collapse of two iron ore tailing dams. Heliyon, 2020, 6, e04778.	1.4	10
879	Additional new species suggest high dictyostelid diversity on Madagascar. Mycologia, 2020, 112, 1026-1042.	0.8	2
881	Thermotogales origin scenario of eukaryogenesis. Journal of Theoretical Biology, 2020, 492, 110192.	0.8	2
882	Identification and subcellular localization of splicing factor arginine/serine-rich 10 in the microsporidian Nosema bombycis. Journal of Invertebrate Pathology, 2020, 174, 107441.	1.5	1

#	Article	IF	CITATIONS
883	Cold climate adaptation is a plausible cause for evolution of multicellular sporulation in Dictyostelia. Scientific Reports, 2020, 10, 8797.	1.6	6
884	Spatiotemporal Patterns in Diversity and Assembly Process of Marine Protist Communities of the Changjiang (Yangtze River) Plume and Its Adjacent Waters. Frontiers in Microbiology, 2020, 11, 579290.	1.5	10
885	Carboniferous smaller Foraminifera: convergences and divergences. Geological Society Special Publication, 2022, 512, 247-326.	0.8	5
886	The Aureochrome Photoreceptor PtAUREO1a Is a Highly Effective Blue Light Switch in Diatoms. IScience, 2020, 23, 101730.	1.9	14
888	NCBI Taxonomy: a comprehensive update on curation, resources and tools. Database: the Journal of Biological Databases and Curation, 2020, 2020, .	1.4	925
889	Severe meningoencephalitis secondary to calvarial invasion of <i>Lagenidium giganteum</i> forma <i>caninum</i> in a dog. Open Veterinary Journal, 2020, 10, 31-38.	0.3	4
890	Increased survival of the honey bee Apis mellifera infected with the microsporidian Nosema ceranae by effective gene silencing. Archives of Insect Biochemistry and Physiology, 2020, 105, e21734.	0.6	9
891	Classifying Metaheuristics: Towards a unified multi-level classification system. Natural Computing, 2022, 21, 155-171.	1.8	35
892	A Highly Conserved Iron-Sulfur Cluster Assembly Machinery between Humans and Amoeba Dictyostelium discoideum: The Characterization of Frataxin. International Journal of Molecular Sciences, 2020, 21, 6821.	1.8	3
893	Land-use change effects on protozoic silicon pools in the Dajiuhu National Wetland Park, China. Geoderma, 2020, 368, 114305.	2.3	10
894	Developmental stages of Amphibiocystidium sp., a parasite from the Italian stream frog (Rana italica). Zoology, 2020, 141, 125813.	0.6	2
895	Structure and Function of Branching Enzymes in Eukaryotes. Trends in Glycoscience and Glycotechnology, 2020, 32, E21-E30.	0.0	5
896	Soil protist communities in burrowing and casting hotspots of different earthworm species. Soil Biology and Biochemistry, 2020, 144, 107774.	4.2	13
897	Functional analysis of the methyltransferase SMYD in the single-cell model organism Tetrahymena thermophila. Marine Life Science and Technology, 2020, 2, 109-122.	1.8	22
898	Evolution and roles of cytokinin genes in angiosperms 2: Do ancient CKXs play housekeeping roles while non-ancient CKXs play regulatory roles?. Horticulture Research, 2020, 7, 29.	2.9	32
899	Microalgae harvesting by fungal-assisted bioflocculation. Reviews in Environmental Science and Biotechnology, 2020, 19, 369-388.	3.9	29
900	New Molecular Approach for the Detection of Kinetoplastida Parasites of Medical and Veterinary Interest. Microorganisms, 2020, 8, 356.	1.6	16
901	Evolution and roles of cytokinin genes in angiosperms 1: Do ancient IPTs play housekeeping while non-ancient IPTs play regulatory roles?. Horticulture Research, 2020, 7, 28.	2.9	27

#	Article	IF	CITATIONS
902	Eukaryotic life without tQCUG: the role of Elongator-dependent tRNA modifications in Dictyostelium discoideum. Nucleic Acids Research, 2020, 48, 7899-7913.	6.5	5
903	Two New Terpenes Isolated from Dictyostelium Cellular Slime Molds. Molecules, 2020, 25, 2895.	1.7	4
904	Orientation of FtsH protease homologs in Trypanosoma brucei inner mitochondrial membrane and its evolutionary implications. Molecular and Biochemical Parasitology, 2020, 238, 111282.	0.5	3
905	Viral RNA Genomes Identified from Marine Macroalgae and a Diatom. Microbes and Environments, 2020, 35, n/a.	0.7	17
906	Morphological description of Opalina obtrigonoidea Metcalf, 1923 (Heterokonta, Opalinea) from Duttaphrynus melanostictus and evaluation of the ITS region as a suitable genetic marker for inter-species identification in Opalina. Parasitology International, 2020, 76, 102103.	0.6	2
908	Landscape of Eukaryotic Transmembrane Beta Barrel Proteins. Journal of Proteome Research, 2020, 19, 1209-1221.	1.8	5
909	Deep Evolutionary History of the Phox and Bem1 (PB1) Domain Across Eukaryotes. Scientific Reports, 2020, 10, 3797.	1.6	13
910	Diversification of CORVET tethers facilitates transport complexity in <i>Tetrahymena thermophila</i> . Journal of Cell Science, 2020, 133, .	1.2	16
911	Comparing Morphological and Molecular Estimates of Species Diversity in the Freshwater Genus <i>Synura</i> (Stramenopiles): A Model for Understanding Diversity of Eukaryotic Microorganisms. Journal of Phycology, 2020, 56, 574-591.	1.0	26
912	Giant virus diversity and host interactions through global metagenomics. Nature, 2020, 578, 432-436.	13.7	207
913	Review on protozoic silica and its role in silicon cycling. Geoderma, 2020, 365, 114224.	2.3	40
914	Global cellulose biomass, horizontal gene transfers and domain fusions drive microbial expansin evolution. New Phytologist, 2020, 226, 921-938.	3.5	19
915	Host Genotypic Effect on Algal Symbiosis Establishment in the Coral Model, the Anemone Exaiptasia diaphana, From the Great Barrier Reef. Frontiers in Marine Science, 2020, 6, .	1.2	23
916	µgreen-db: a reference database for the 23S rRNA gene of eukaryotic plastids and cyanobacteria. Scientific Reports, 2020, 10, 5915.	1.6	17
917	A review of the systematics, species identification and diagnostics of the Trypanosomatidae using the maxicircle kinetoplast DNA: from past to present. International Journal for Parasitology, 2020, 50, 449-460.	1.3	5
918	Remarkably preserved cysts of the extinct synurophyte, Mallomonas ampla, uncovered from a 48 Ma freshwater Eocene lake. Scientific Reports, 2020, 10, 5204.	1.6	1
919	Checklist of parasites for Pontoâ€Caspian gobies (Actinopterygii: Gobiidae) in their native and nonâ€native ranges. Journal of Applied Ichthyology, 2020, 36, 472-500.	0.3	11
920	Specificities of the plant mitochondrial translation apparatus. Mitochondrion, 2020, 53, 30-37.	1.6	10

		CITATION RE	PORT	
#	Article		IF	CITATIONS
921	The Syntrophy hypothesis for the origin of eukaryotes revisited. Nature Microbiology, 2	2020, 5, 655-667.	5.9	104
922	Comparative analyses of the V4 and V9 regions of 18S rDNA for the extant eukaryotic the Illumina platform. Scientific Reports, 2020, 10, 6519.	community using	1.6	51
924	The Ecology and Evolution of Amoeba-Bacterium Interactions. Applied and Environmer Microbiology, 2021, 87, .	Ital	1.4	42
925	Contaminated water confirmed as source of infection by bioassay in an outbreak of to South Brazil. Transboundary and Emerging Diseases, 2021, 68, 767-772.	xoplasmosis in	1.3	25
926	Metabolomic approach of the antiprotozoal activity of medicinal Piper species used in Amazon. Journal of Ethnopharmacology, 2021, 264, 113262.	Peruvian	2.0	10
927	Proteome-Scale Detection of Differential Conservation Patterns at Protein and Subprowith BLUR. Genome Biology and Evolution, 2021, 13, .	tein Levels	1.1	3
928	High Sequence Divergence but Limited Architectural Rearrangements in Organelle Gen Cyanophora (Glaucophyta) Species. Journal of Eukaryotic Microbiology, 2021, 68, e12		0.8	0
929	Molecular detection and characterization of Theileria sp. from hedgehogs (Paraechinu in Saudi Arabia. Letters in Applied Microbiology, 2021, 72, 476-483.	s aethiopicus)	1.0	2
930	Application of low dosage of copper oxide and zinc oxide nanoparticles boosts bacteria communities in soil. Science of the Total Environment, 2021, 757, 143807.	al and fungal	3.9	26
931	Prokaryotic and microeukaryotic communities in an experimental rice plantation under of pesticides. Environmental Science and Pollution Research, 2021, 28, 2328-2341.	long-term use	2.7	10
932	Unraveling the sugar code: the role of microbial extracellular glycans in plant–microb Journal of Experimental Botany, 2021, 72, 15-35.	e interactions.	2.4	37
934	Novel biosensors for detection of the parasite in food. , 2021, , 59-77.			1
935	Morphology and classification. , 2021, , 59-97.			0
936	tRNA 3′ shortening by LCCR4 as a response to stress in Trypanosoma brucei. Nuclei 2021, 49, 1647-1661.	c Acids Research,	6.5	9
937	Haemogregarines and Criteria for Identification. Animals, 2021, 11, 170.		1.0	9
938	Gap Analysis of Threatened, Rare, and Under-Represented Species in Bhutan. , 2021, , 2	199-278.		0
939	Phylogenomic fingerprinting of tempo and functions of horizontal gene transfer withir ochrophytes. Proceedings of the National Academy of Sciences of the United States of 118, .		3.3	37
940	Biodiversity of the Coccidia (Apicomplexa: Conoidasida) in vertebrates: what we know, not know, and what needs to be done. Folia Parasitologica, 2021, 68, .	what we do	0.7	7

	Сітатіо	n Report	
# 941	ARTICLE Nomenclature: how do we designate NPP taxa?. Geological Society Special Publication, 2021, 511, 77-89.	IF 0.8	CITATIONS
941	Nomenciature. Now do we designate NPP taxa:. Geological Society Special Publication, 2021, 311, 77-05.	0.8	0
942	How Does Sphagnum Growing Affect Testate Amoeba Communities and Corresponding Protozoic Si Pools? Results from Field Analyses in SW China. Microbial Ecology, 2021, 82, 459-469.	1.4	7
943	In Silico Identification and Functional Characterization of Conserved miRNAs in the Genome of <i>Cryptosporidium parvum</i> . Bioinformatics and Biology Insights, 2021, 15, 117793222110276.	1.0	4
944	Algae for global sustainability?. Advances in Botanical Research, 2021, , 145-212.	0.5	9
945	Arctic biodiversity amidst looming climate apocalypse: current status and way forward. , 2021, , 213-255.		0
947	Ediacaran metazoan fossils with siliceous skeletons from the Digermulen Peninsula of Arctic Norway. Journal of Paleontology, 2021, 95, 440-475.	0.5	4
948	Peptidylarginine Deiminase (PAD) and Post-Translational Protein Deimination—Novel Insights into Alveolata Metabolism, Epigenetic Regulation and Host–Pathogen Interactions. Biology, 2021, 10, 177.	1.3	4
949	RNA Viruses in Aquatic Unicellular Eukaryotes. Viruses, 2021, 13, 362.	1.5	25
950	Influence of Zwitterionic Buffer Effects with Thermal Modification Treatments of Wood on Symbiotic Protists in Reticulitermes grassei Clément. Insects, 2021, 12, 139.	1.0	5
951	Silicon Cycling in Soils Revisited. Plants, 2021, 10, 295.	1.6	105
952	The Hulks and the Deadpools of the Cytokinin Universe: A Dual Strategy for Cytokinin Production, Translocation, and Signal Transduction. Biomolecules, 2021, 11, 209.	1.8	35
953	Identification and localization of Nup170 in the microsporidian Nosema bombycis. Parasitology Research, 2021, 120, 2125-2134.	0.6	1
954	Structure and dynamics of the periphytic ciliate community under different hydrological conditions in a Danubian floodplain lake. Limnologica, 2021, 87, 125847.	0.7	5
955	Protists as main indicators and determinants of plant performance. Microbiome, 2021, 9, 64.	4.9	71
957	Protocol optimization for the detection of Trypanosoma cruzi DNA in açai (Euterpe oleraceae) pulp. Acta Amazonica, 2021, 51, 79-84.	0.3	1
958	Gene Duplications Trace Mitochondria to the Onset of Eukaryote Complexity. Genome Biology and Evolution, 2021, 13, .	1.1	24
960	<i>Olisthodiscus</i> represents a new class of Ochrophyta. Journal of Phycology, 2021, 57, 1094-1118.	1.0	10
961	Characterization of the RNA-interference pathway as a tool for reverse genetic analysis in the nascent phototrophic endosymbiosis, <i>Paramecium bursaria</i> . Royal Society Open Science, 2021, 8, 210140.	1.1	6

		CITATION RE	EPORT	
#	Article		IF	CITATIONS
962	Cadherins in early neural development. Cellular and Molecular Life Sciences, 2021, 78,	4435-4450.	2.4	13
963	The curious consistency of carbon biosignatures over billions of years of Earth-life coev ISME Journal, 2021, 15, 2183-2194.	volution.	4.4	26
964	Microeukaryotic and Prokaryotic Diversity of Anchialine Caves from Eastern Adriatic Se Microbial Ecology, 2022, 83, 257-270.	a Islands.	1.4	9
965	The Unicellular Red Alga <i>Cyanidioschyzon merolae—</i> The Simplest Model of a P Eukaryote. Plant and Cell Physiology, 2021, 62, 926-941.	hotosynthetic	1.5	24
966	Eco-Evolutionary Perspectives on Mixoplankton. Frontiers in Marine Science, 2021, 8, .		1.2	10
968	De Novo Transcriptome Meta-Assembly of the Mixotrophic Freshwater Microalga Eugle Genes, 2021, 12, 842.	ena gracilis.	1.0	9
969	Analysis of diverse eukaryotes suggests the existence of an ancestral mitochondrial ap from the bacterial type II secretion system. Nature Communications, 2021, 12, 2947.	paratus derived	5.8	19
970	Sequential production of gametes during meiosis in trypanosomes. Communications E 555.	Biology, 2021, 4,	2.0	18
971	Giardiavirus: an update. Parasitology Research, 2021, 120, 1943-1948.		0.6	6
972	Structural basis of ubiquitination mediated by protein splicing in early Eukarya. Biochir Biophysica Acta - General Subjects, 2021, 1865, 129844.	nica Et	1.1	2
973	Large-scale phylogenomic analysis provides new insights into the phylogeny of the clas Oligohymenophorea (Protista, Ciliophora) with establishment of a new subclass Uroce subcl. Molecular Phylogenetics and Evolution, 2021, 159, 107112.		1.2	23
974	Direct visualization of epithelial microvilli biogenesis. Current Biology, 2021, 31, 2561-	2575.e6.	1.8	28
975	Sialic Acids as Receptors for Pathogens. Biomolecules, 2021, 11, 831.		1.8	27
976	Low shifts in salinity determined assembly processes and network stability of microeul plankton communities in a subtropical urban reservoir. Microbiome, 2021, 9, 128.	karyotic	4.9	191
977	The sediment reworking foraminiferan Ammonia cf. aomoriensis is a sediment destabil from an experiment with artificial removal of the pseudopods. Regional Studies in Mari 2021, 45, 101814.		0.4	1
978	Phagotrophic protists (protozoa) in Antarctic terrestrial ecosystems: diversity, distribu and best research practices. Polar Biology, 2021, 44, 1467-1484.	tion, ecology,	0.5	1
979	Clypifer cribrifer gen. nov., sp. nov. (Clypiferidae fam. nov., Pterocystida, Centroplasthe notes on evolution of centrohelid siliceous coverings. International Journal of Systema Evolutionary Microbiology, 2021, 71, .		0.8	6
980	Dumpster diving for diatom plastid 16S rRNA genes. PeerJ, 2021, 9, e11576.		0.9	3

ARTICLE

IF CITATIONS

A new integrated morpho- and molecular systematic classification of Cenozoic radiolarians (Class) Tj ETQq0 0 0 rgBT $_{0.2}^{1}$ Overlock 10 Tf 50

983	The longâ€ŧime orphan protist <i>Meringosphaeramediterranea</i> Lohmann, 1902 [1903] is a centrohelid heliozoan. Journal of Eukaryotic Microbiology, 2021, 68, e12860.	0.8	4
984	The First Discovery of Colonial Radiolarians in Jurassic Sediments (Arctic Zone of Siberia). Paleontological Journal, 2021, 55, 355-361.	0.2	0
985	ensembleTax: an R package for determinations of ensemble taxonomic assignments of phylogenetically-informative marker gene sequences. PeerJ, 2021, 9, e11865.	0.9	3
986	New gregarine species (Apicomplexa) from tunicates show an evolutionary history of host switching and suggest a problem with the systematics of Lankesteria and Lecudina. Journal of Invertebrate Pathology, 2021, 183, 107622.	1.5	2
987	Origin and Early Evolution of the Eukaryotic Cell. Annual Review of Microbiology, 2021, 75, 631-647.	2.9	28
988	Comparative analysis of freshwater phytoplankton communities in two lakes of Burabay National Park using morphological and molecular approaches. Scientific Reports, 2021, 11, 16130.	1.6	12
989	Winogradsky columns as a strategy to study typically rare microbial eukaryotes. European Journal of Protistology, 2021, 80, 125807.	0.5	4
990	An update on cell division of Giardia duodenalis trophozoites. Microbiological Research, 2021, 250, 126807.	2.5	7
991	Histopathological description of an emerging disease in Norwegian salmonid aquaculture caused by an xâ€cell parasite. Journal of Fish Diseases, 2022, 45, 213-217.	0.9	2
992	The specificity of Babesia-tick vector interactions: recent advances and pitfalls in molecular and field studies. Parasites and Vectors, 2021, 14, 507.	1.0	15
993	Bioprospecting of thraustochytrids for omega-3 fatty acids: A sustainable approach to reduce dependency on animal sources. Trends in Food Science and Technology, 2021, 115, 433-444.	7.8	38
994	Early-diverging fungal phyla: taxonomy, species concept, ecology, distribution, anthropogenic impact, and novel phylogenetic proposals. Fungal Diversity, 2021, 109, 59-98.	4.7	35
995	Gamete expression of TALE class HD genes activates the diploid sporophyte program in Marchantia polymorpha. ELife, 2021, 10, .	2.8	35
996	Soil aggregate isolation method affects interpretation of protistan community. Soil Biology and Biochemistry, 2021, 161, 108388.	4.2	8
997	Co-occurrence patterns and assembly processes of microeukaryotic communities in a semi-enclosed aquaculture bay. Continental Shelf Research, 2021, 228, 104550.	0.9	6
998	Tracking microeukaryotic footprint in a peri-urban watershed, China through machine-learning approaches. Science of the Total Environment, 2022, 806, 150401.	3.9	15
999	The phylogeny and phylogenetically based classification of myxomycetes. , 2022, , 97-124.		3

#	ARTICLE Phosphagen kinases from five groups of eukaryotic protists (Choanomonada, Alveolate,) Tj ETQq0 0 0 rgBT /Ove	IF rlock 10 ⁻¹	CITATIONS Tf 50 752 Td (
1000	with metazoan enzymes. Comparative Biochemistry and Physiology - B Biochemistry and Molecular Biology, 2022, 257, 110663.	0.7	2
1001	Taxonomy and systematics: current knowledge and approaches on the taxonomic treatment of Myxomycetes: updated version. , 2022, , 269-324.		1
1002	Key drivers influencing the colonization of periphytic ciliates and their functional role in hydrologically dynamic floodplain lake ecosystem. Knowledge and Management of Aquatic Ecosystems, 2021, , 33.	0.5	3
1003	LIM homeodomain proteins and associated partners: Then and now. Current Topics in Developmental Biology, 2021, 145, 113-166.	1.0	15
1008	Diatom Classifications: What Purpose Do They Serve?. , 2020, , 11-24.		6
1009	Protosteloid Amoebae (Protosteliida, Protosporangiida, Cavosteliida, Schizoplasmodiida,) Tj ETQq1 1 0.784314 r	gBT /Ove	rlock 10 Tf 50
1010	Dictyostelia. , 2017, , 1-45.		1
1011	Apicomplexa. , 2016, , 1-58.		20
1012	Hyphochytriomycota and Oomycota. , 2016, , 1-71.		9
1013	Retortamonadida (with Notes on Carpediemonas-Like Organisms and Caviomonadidae). , 2016, , 1-32.		3
1014	Eustigmatophyceae. , 2016, , 1-39.		5
1015	Protist Diversity and Eukaryote Phylogeny. , 2017, , 1-21.		4
1016	Parabasalia. , 2016, , 1-44.		9
1017	Phytoplankton Responses to Marine Climate Change – An Introduction. , 2018, , 55-71.		16
1018	Gregarines. , 2015, , 1-47.		6
1019	An Evolutionary Perspective on the Plant Hormone Ethylene. , 2015, , 109-134.		4
1020	Timing the Origins of Multicellular Eukaryotes Through Phylogenomics and Relaxed Molecular Clock Analyses. Advances in Marine Genomics, 2015, , 3-29.	1.2	19
1021	Peptides from diatoms and grasses harness phosphate ion binding to silica to help regulate biomaterial structure. Acta Biomaterialia, 2020, 112, 286-297.	4.1	6

#	Article	IF	CITATIONS
1022	Particles, protists, and zooplankton in glacier-influenced coastal Svalbard waters. Estuarine, Coastal and Shelf Science, 2020, 242, 106842.	0.9	10
1023	Mutual antagonism between Hippo signaling and cyclin E drives intracellular pattern formation. Journal of Cell Biology, 2020, 219, .	2.3	7
1024	The small subunit rRNA gene sequence of the chonotrich Chilodochona carcini Jankowski, 1973 confirms chonotrichs as a dysteriid-derived clade (Phyllopharyngea, Ciliophora). International Journal of Systematic and Evolutionary Microbiology, 2016, 66, 2959-2964.	0.8	10
1025	Identification of Pythium insidiosum complex by matrix-assisted laser desorption ionization-time of flight mass spectrometry. Journal of Medical Microbiology, 2019, 68, 574-584.	0.7	21
1036	Classification of Reynolds phytoplankton functional groups using individual traits and machine learning techniques. Freshwater Biology, 2017, 62, 1681-1692.	1.2	55
1037	Exclusive Gut Flagellates of Serritermitidae Suggest a Major Transfaunation Event in Lower Termites: Description of <i>Heliconympha glossotermitis</i> gen. nov. spec. nov Journal of Eukaryotic Microbiology, 2018, 65, 77-92.	0.8	29
1038	<i>Parvularia atlantis</i> gen. et sp. nov., a Nucleariid Filose Amoeba (Holomycota, Opisthokonta). Journal of Eukaryotic Microbiology, 2018, 65, 170-179.	0.8	21
1039	Assessing SSU rDNA Barcodes in Foraminifera: A Case Study using Bolivina quadrata. Journal of Eukaryotic Microbiology, 2018, 65, 220-235.	0.8	6
1041	Taxonomy and Classification of Human Parasitic Protozoa and Helminths. , 0, , 2282-2292.		1
1042	The Protists in Soil—A Token of Untold Eukaryotic Diversity. , 2019, , 125-140.		15
1043	Distribution of Myxomycetes Among the Microhabitats Available for these Organisms in Tropical Forests. , 2014, , 134-151.		7
1045	Recapitulating phylogenies using k-mers: from trees to networks. F1000Research, 2016, 5, 2789.	0.8	22
1046	Draft genome assembly and transcriptome sequencing of the golden algae Hydrurus foetidus (Chrysophyceae). F1000Research, 2019, 8, 401.	0.8	8
1047	Recent advances in understanding mitochondrial genome diversity. F1000Research, 2020, 9, 270.	0.8	63
1048	The ubiquitous and ancient ER membrane protein complex (EMC): tether or not?. F1000Research, 2015, 4, 624.	0.8	75
1049	The ubiquitous and ancient ER membrane protein complex (EMC): tether or not?. F1000Research, 2015, 4, 624.	0.8	63
1050	From simple to supercomplex: mitochondrial genomes of euglenozoan protists. F1000Research, 0, 5, 392.	0.8	5
1051	From simple to supercomplex: mitochondrial genomes of euglenozoan protists. F1000Research, 2016, 5, 392.	0.8	12

#	Article	IF	CITATIONS
1052	Old Lineages in a New Ecosystem: Diversification of Arcellinid Amoebae (Amoebozoa) and Peatland Mosses. PLoS ONE, 2014, 9, e95238.	1.1	15
1053	Mdb1, a Fission Yeast Homolog of Human MDC1, Modulates DNA Damage Response and Mitotic Spindle Function. PLoS ONE, 2014, 9, e97028.	1.1	23
1054	Heterogeneous Occupancy and Density Estimates of the Pathogenic Fungus Batrachochytrium dendrobatidis in Waters of North America. PLoS ONE, 2014, 9, e106790.	1.1	75
1055	Molecular Profiling of the Phytophthora plurivora Secretome: A Step towards Understanding the Cross-Talk between Plant Pathogenic Oomycetes and Their Hosts. PLoS ONE, 2014, 9, e112317.	1.1	16
1056	Protococcidian Eleutheroschizon duboscqi, an Unusual Apicomplexan Interconnecting Gregarines and Cryptosporidia. PLoS ONE, 2015, 10, e0125063.	1.1	17
1057	"Candidatus Fokinia solitariaâ€, a Novel "Stand-Alone―Symbiotic Lineage of Midichloriaceae (Rickettsiales). PLoS ONE, 2016, 11, e0145743.	1.1	44
1058	RuBisCO in Non-Photosynthetic Alga Euglena longa: Divergent Features, Transcriptomic Analysis and Regulation of Complex Formation. PLoS ONE, 2016, 11, e0158790.	1.1	31
1059	Multispecies reconstructions uncover widespread conservation, and lineage-specific elaborations in eukaryotic mRNA metabolism. PLoS ONE, 2018, 13, e0192633.	1.1	20
1060	An Interactome-Centered Protein Discovery Approach Reveals Novel Components Involved in Mitosome Function and Homeostasis in Giardia lamblia. PLoS Pathogens, 2016, 12, e1006036.	2.1	22
1061	Specialising the parasite nucleus: Pores, lamins, chromatin, and diversity. PLoS Pathogens, 2017, 13, e1006170.	2.1	11
1062	Morphological switch to a resistant subpopulation in response to viral infection in the bloom-forming coccolithophore Emiliania huxleyi. PLoS Pathogens, 2017, 13, e1006775.	2.1	29
1064	Four different <i>Phytophthora</i> species that are able to infect Scots pine seedlings in laboratory conditions. Folia Forestalia Polonica, Series A, 2016, 58, 123-130.	0.1	2
1065	The prospects and perspectives of the phylogenetic system of myxomycetes (Myxogastrea). Ukrainian Botanical Journal, 2015, 72, 147-155.	0.1	3
1066	Development of Knowledge of the Taxonomy and Phylogeny of Living Organisms for Future Biology Teachers. The Advanced Science Journal, 2015, 2015, 75-82.	0.0	1
1067	Prokaryotic ancestry and gene fusion of a dual localized peroxiredoxin in malaria parasites. Microbial Cell, 2015, 2, 5-13.	1.4	9
1068	The Coccidia (Apicomplexa) of the Archosauria (Crocodylia: Eusuchia) of the World. Journal of Parasitology, 2020, 106, 90.	0.3	5
1069	RNA interference in formation of the somatic genome of ciliates Paramecium and Tetrahymena. Ecological Genetics, 2018, 16, 5-22.	0.1	4
1070	Evolutionary history of the group formerly known as protists using a phylogenomics approach. Revista De La Academia Colombiana De Ciencias Exactas, Fisicas Y Naturales, 2016, 40, 147.	0.0	1

_	
Citation	
CHAHON	REPORT

#	Article	IF	CITATIONS
1072	DISENTANGLING THE TAXONOMIC STRUCTURE OF THE LEPIDODERMA CHAILLETII-CARESTIANUM SPECIES COMPLEX (MYXOGASTRIA, AMOEBOZOA): GENETIC AND MORPHOLOGICAL ASPECTS. Protistology, 2016, 10, .	0.0	16
1073	Heterotrophic flagellates and centrohelid heliozoa from littoral and supralittoral zones of the Black Sea (the Southern part of the Crimea). Protistology, 2017, 11, .	0.0	4
1074	Free-living heterotrophic flagellates from bays of Sevastopol (the Black Sea littoral). Protistology, 2018, 12, .	0.0	4
1075	Gene and Chromosomal Copy Number Variations as an Adaptive Mechanism Towards a Parasitic Lifestyle in Trypanosomatids. Current Genomics, 2018, 19, 87-97.	0.7	44
1076	Analysis of DNA Exchange Using Thymidine Analogs (ADExTA) in Trypanosoma cruzi. Bio-protocol, 2018, 8, e3125.	0.2	4
1077	Wind drives microbial eukaryote communities in a temperate closed lagoon. Aquatic Microbial Ecology, 2017, 78, 187-200.	0.9	9
1078	Neurotropic mesomycetozoean-like infection in larvae of the southern toad Anaxyrus terrestris in Florida, USA. Diseases of Aquatic Organisms, 2015, 113, 157-162.	0.5	2
1079	Diversity and microhabitat associations of Labyrinthula spp. in the Indian River Lagoon System. Diseases of Aquatic Organisms, 2020, 137, 145-157.	0.5	6
1080	Seasonality in the vertical flux and species composition of Radiolaria Polycystina (Protista): patterns and drivers. Marine Ecology - Progress Series, 2017, 578, 51-72.	0.9	2
1081	Checklist of Sphagnum-dwelling testate amoebae in Bulgaria. Biodiversity Data Journal, 2018, 6, e25295.	0.4	5
1082	Morphologic and molecular data help adopting the insect-pathogenic nephridiophagids (Nephridiophagidae) among the early diverging fungal lineages, close to the Chytridiomycota. MycoKeys, 0, 25, 31-50.	0.8	14
1083	Evolutionary analysis of Slc11 mechanism of proton-coupled metal-ion transmembrane import. AIMS Biophysics, 2016, 3, 286-318.	0.3	6
1084	Check-list of microscopic protosteloid amoebae from the Southwest of Europe. Anales Del Jardin Botanico De Madrid, 2012, 69, 217-236.	0.2	1
1085	Preservice biology teachers' classificatory thinking about plants. Recherches En Didactiques Des Sciences Et Des Technologies, 2017, , 57-92.	0.1	2
1090	Preliminary studies of fungi in the Biebrza National Park (NE Poland). Part III. Micromycetes – new data. Acta Mycologica, 2016, 50, .	0.3	2
1091	Punctuated evolution and transitional hybrid network in an ancestral cell cycle of fungi. ELife, 2016, 5, .	2.8	52
1092	A diverse host thrombospondin-type-1 repeat protein repertoire promotes symbiont colonization during establishment of cnidarian-dinoflagellate symbiosis. ELife, 2017, 6, .	2.8	44
1093	Comprehensive transcriptome analysis provides new insights into nutritional strategies and phylogenetic relationships of chrysophytes. PeerJ, 2017, 5, e2832.	0.9	38

		CHAHON REPORT		
#	ARTICLE	IF	CITATIONS	
1094	Microbial communities in the reef water at Kham Island, lower Gulf of Thailand. PeerJ, 2017, 5, e3625.	0.9	9	
1095	Diatom Resting Stages in Surface Sediments: A Pilot Study Comparing Next Generation Sequencing and Serial Dilution Cultures. Cryptogamie, Algologie, 2017, 38, 31-46.	0.3	28	
1096	On the conservation of fungi. Agricultural Science Euro-North-East, 2021, 22, 641-660.	0.2	1	
1097	Ecological and evolutionary diversification of sulphated polysaccharides in diverse photosynthetic lineages: A review. Carbohydrate Polymers, 2022, 277, 118764.	5.1	8	
1098	Sexual Development in Non-Human Parasitic Apicomplexa: Just Biology or Targets for Control?. Animals, 2021, 11, 2891.	1.0	9	
1099	Cellular and Molecular Targets of Nucleotide-Tagged Trithiolato-Bridged Arene Ruthenium Complexes in the Protozoan Parasites Toxoplasma gondii and Trypanosoma brucei. International Journal of Molecular Sciences, 2021, 22, 10787.	1.8	13	
1100	Modern Benthic Foraminiferal Diversity Along the Fjords of Svalbard Archipelago: Diversity Evaluation. Thalassas, 2022, 38, 647-664.	0.1	2	
1101	Expression and production of thermophilic alginate lyases in Bacillus and direct application of culture supernatant for seaweed saccharification. Algal Research, 2021, 60, 102512.	2.4	3	
1102	Introduction to Cryptosporidium: The Parasite and the Disease. SpringerBriefs in Food, Health and Nutrition, 2014, , 1-10.	0.5	0	
1103	Preliminary Study on Phytoplankton Distribution Changes Monitoring for the Intensive Study Area of the Ariake Sea, Japan Based on Remote Sensing Satellite Data. International Journal of Advanced Research in Artificial Intelligence, 2014, 3, .	0.2	0	
1105	Free-living Amoebae (FLA), Introduction. , 2015, , 1-3.		0	
1106	ls cryptosporidiosis an underestimated disease in cats?. Archivos De Medicina Veterinaria, 2015, 47, 1-6.	0.2	0	
1107	Taxonomy and Classification of Fungi. , 0, , 1932-1943.		0	
1109	Endosymbiosis in <i>Paramecium bursaria</i> . Hikaku Seiri Seikagaku(Comparative Physiology and) Tj ETQq1 1	0.784314	rg&T /Overlo	
1110	Toward the establishment of glaucophyte taxonomic system based on the threedimensional recognition of ultrastructures. Plant Morphology, 2016, 28, 49-54.	0.1	0	
1111	Heterolobosea. , 2016, , 1-42.		1	
1112	Free-Living Amoebae (FLA), Introduction. , 2016, , 1074-1076.		0	
1113	Centrohelida and Other Heliozoan-Like Protists. , 2016, , 1-17.		0	

#	Article	IF	CITATIONS
1114	Amoebozoan Lobose Amoebae (Tubulinea, Flabellinea, and Others). , 2016, , 1-31.		2
1115	Gregarines. , 2016, , 1142-1188.		0
1116	Myxomycetes. , 2016, , 1-27.		2
1117	Radiolaria and Phaeodaria. , 2016, , 1-33.		0
1118	Zygnematophyta. , 2016, , 1-30.		1
1121	Charophytic algae: issues of evolution and phylogeny. Ukrainian Botanical Journal, 2016, 73, 163-177.	0.1	1
1122	Invertebrate Collection Donated by Professor Dr. Ion Cantacuzino to "Grigore Antipa―National Museum of Natural History from Bucharest. Travaux Du Museum National D'Histoire Naturelle Grigore Antipa, 2016, 59, 7-30.	0.1	0
1123	Recapitulating phylogenies using k-mers: from trees to networks. F1000Research, 2016, 5, 2789.	0.8	13
1124	Zygnematophyta. , 2017, , 1-29.		0
1125	Paramyxida. , 2017, , 1-18.		0
1126	Radiolaria and Phaeodaria. , 2017, , 1-33.		2
1132	Control of Flagellum Length by a Grow-and-Lock Model. SSRN Electronic Journal, 0, , .	0.4	0
1133	Root Gall Formation, Resting Spore Isolation and High Molecular Weight DNA Extraction of Plasmodiophora brassicae. Bio-protocol, 2018, 8, .	0.2	1
1136	Milky-White Bloom in Lake Oshima-ohnuma Caused by an Algae-lysing Protist, <i>Asterocaelum</i> sp Japanese Journal of Limnology, 2018, 79, 109-117.	0.1	4
1137	Symbiosis in Eukaryotic Cell Evolution. , 2018, , 119-148.		0
1140	Phytopathogenic fungi of the Ukrainian Roztochya (collected in the spring of 2016 and 2017 years). Proceedings of the State Natural History Museum, 2018, , 125-134.	0.2	0
1143	Live protist curation at the Scottish Association for Marine Science, 1884–2017. Archives of Natural History, 2018, 45, 267-282.	0.0	1
1144	Something for Everyone: A Review of "The Biology and Identification of the Coccidia (Apicomplexa) of Carnivores of the World― American Midland Naturalist, 2019, 181, 143.	0.2	1

#	Article	IF	CITATIONS
1145	Asgardarchaeota – A Novel Prokaryotic Group Discovered in Aquatic Sediments that Might Shed Light on the Origin and Early Evolution of Eukaryotes , 2019, , .		1
1146	Draft genome assembly and transcriptome sequencing of the golden algae Hydrurus foetidus (Chrysophyceae). F1000Research, 2019, 8, 401.	0.8	3
1150	Amoebozoa Luhe, 1913 in the water bodies of the Vinnytsia region. Bulletin of Taras Shevchenko National University of Kyiv Series Biology, 2020, 82, 48-52.	0.1	0
1151	16 Biotechnology of Marine Fungi: New Workhorses and Applications. , 2020, , 399-412.		1
1153	Accumulation of Tetrahymena pyriformis on Interfaces. Micromachines, 2021, 12, 1339.	1.4	6
1154	The World of Algae Reveals a Broad Variety of Cryptochrome Properties and Functions. Frontiers in Plant Science, 2021, 12, 766509.	1.7	20
1156	Spezielle Zoologie Teil A: "Protista" und Metazoa. , 2020, , 37-71.		0
1159	Structure and Function of Branching Enzymes in Eukaryotes. Trends in Glycoscience and Glycotechnology, 2020, 32, J19-J28.	0.0	Ο
1160	A New Testate Amoeba, Matsakision ogawaraensis sp. nov. (Silicofilosea: Incertae sedis Euglyphida) from Lake Shore Sand of Northern Japan. Acta Protozoologica, 2020, 59, 99-105.	0.5	1
1162	Ecology and Distribution. , 2020, , 13-22.		0
1164	Sar S. M. Adl et al. 2012 [A. G. B. Simpson and M. Dunthorn], converted clade name. , 2020, , 51-54.		0
1166	Eukarya R. Creti et al. 1991 [A. G. B. Simpson], converted clade name. , 2020, , 29-32.		Ο
1167	Fungi A. Engler and K. Prantl 1900: iii [D. S. Hibbett, M. Blackwell, T. James, J. W. Spatafora, J. W. Taylor, and R. Vilgalys], converted clade name. , 2020, , 109-114.		0
1168	<i>Txikispora philomaios</i> n. sp., n. g., a microâ€eukaryotic pathogen of amphipods, reveals parasitism and hidden diversity in Class Filasterea. Journal of Eukaryotic Microbiology, 2022, 69, e12875.	0.8	6
1169	Distribution of naked amoebae in the soils of the steppe zone of Ukraine. Visnyk L'vivs'koho Universytetu Seriia Biolohichna, 2020, , 159-166.	0.0	0
1171	Detection and Control of Fungal Outbreaks. Mycopathologia, 2020, 185, 741-745.	1.3	4
1174	Metagenomics approach for Polymyxa betae genome assembly enables comparative analysis towards deciphering the intracellular parasitic lifestyle of the plasmodiophorids. Genomics, 2022, 114, 9-22.	1.3	4
1175	Actualización taxonómica de las algas pardas (Phaeophyceae, Ochrophyta) marinas bentónicas del Atlántico mexicano. Acta Botanica Mexicana, 2021, , .	0.1	4

	Сіта	CITATION REPORT		
# 1176	ARTICLE Global marine phytoplankton revealed by the Tara Oceans expedition. , 2022, , 531-561.	IF	CITATIONS 2	
1177	Soil protists: An untapped microbial resource of agriculture and environmental importance. Pedosphere, 2022, 32, 184-197.	2.1	23	
1178	Plant and animal two-pore channels. , 2022, , 247-267.		0	
1179	Basal Parasitic Fungi in Marine Food Webs—A Mystery Yet to Unravel. Journal of Fungi (Basel,) Tj ETQq	1 1 0.784314 rgBT 1.5	/Qverlock 1	
1180	Horizontal and vertical distribution of Gambierdiscus spp. (Dinophyceae) including novel phylotypes in Japan identified by 18S rDNA metabarcoding. Harmful Algae, 2022, 111, 102163.	2.2	7	
1181	Experimental Pythium aphanidermatum Infection in Rabbits. Journal of Comparative Pathology, 2022, 190, 30-35.	0.1	0	
1183	A snapshot of parasites in tropical and subtropical freshwater wetlands: modest attention for major players. , 2022, , 417-485.		2	
1184	DBHR: a collection of databases relevant to human research. Future Science OA, 2022, 8, FSO780.	0.9	3	
1185	Forensic significance in the cases of drowning deaths: An elaborative study. IP International Journal of Forensic Medicine and Toxicological Sciences, 2022, 6, 122-126.	0.0	1	
1187	Oxygen gradients shape the unique structure of picoeukaryotic communities in the Bay of Bengal. Science of the Total Environment, 2022, 814, 152862.	3.9	7	
1188	Interactions between soil protists and pollutants: An unsolved puzzle. Journal of Hazardous Materials, 2022, 429, 128297.	6.5	21	
1189	Fractional 2′-O-methylation in the ribosomal RNA of Dictyostelium discoideum supports ribosome heterogeneity in Amoebozoa. Scientific Reports, 2022, 12, 1952.	1.6	4	
1190	Sequence Analysis and Comparison of TCTP Proteins from Human Protozoan Parasites. Acta Parasitologica, 2022, 67, 1024-1031.	0.4	2	
1191	A putative homologue regulates lipid droplet biogenesis via PAH1 in. Journal of Biosciences, 2018, 43, 693-706.	0.5	2	
1193	Supergroups of eukaryotes through a biotechnologist's look. The system of eukaryotes and the need for a taxonomic/biotechnological interface. Pharmacy Formulas, 0, , .	d 0.2	0	
1194	Evolution of circadian clocks along the green lineage. Plant Physiology, 2022, 190, 924-937.	2.3	15	
1195	Free-living protozoa (ciliophora, excavata and amoebozoa) in two water sources for human supply in the municipality of Blumenau, SC. Ciência E Natura, 0, 44, e4.	0.0	0	
1196	Protozoan phagotrophy from predators to parasites: An overview of the enigmatic cytostome ytopharynx complex of <i>Trypanosoma cruzi</i> . Journal of Eukaryotic Microbiology, 2022, 69, e12896.	0.8	4	

#	Article	IF	CITATIONS
1197	Can Aggregate-Associated Organisms Influence the Fouling in a SWRO Desalination Plant?. Microorganisms, 2022, 10, 682.	1.6	1
1198	<i>Paramecium</i> epigenetics in development and proliferation. Journal of Eukaryotic Microbiology, 2022, 69, e12914.	0.8	7
1199	Morphological and phylogenetic investigations shed light on evolutionary relationships of the enigmatic genus Copemetopus (Ciliophora, Alveolata), with the proposal of Copemetopus verae sp. nov European Journal of Protistology, 2022, 83, 125878.	0.5	10
1200	Monophyly of diverse Bigyromonadea and their impact on phylogenomic relationships within stramenopiles. Molecular Phylogenetics and Evolution, 2022, 171, 107468.	1.2	7
1201	Ciliary transition zone evolution and the root of the eukaryote tree: implications for opisthokont origin and classification of kingdoms Protozoa, Plantae, and Fungi. Protoplasma, 2022, 259, 487-593.	1.0	9
1202	High throughput single-cell genome sequencing gives insights into the generation and evolution of mosaic aneuploidy in <i>Leishmania donovani</i> . Nucleic Acids Research, 2022, 50, 293-305.	6.5	14
1204	The Myxobiota of the Å a giewnicki Forest in Å $ ilde{A}^3$ dÅ $^{ m o}$ (Central Poland). Acta Mycologica, 0, 56, .	0.3	0
1205	Above―and belowâ€ground biodiversity responses to the prolonged flood pulse in centralâ€western Amazonia, Brazil. Environmental DNA, 2022, 4, 533-548.	3.1	1
1206	Geographic distribution of Pythium insidiosum infections in the United States. Journal of the American Veterinary Medical Association, 2022, 260, 530-534.	0.2	6
1207	Inferring Species Compositions of Complex Fungal Communities from Long- and Short-Read Sequence Data. MBio, 2022, 13, e0244421.	1.8	2
1228	The Lytic Cycle of Human Apicomplexan Parasites. , 2022, , .		0
1230	Species and morphotypes complexes of naked amoebae in several types of soils of Ukraine. , 2022, 77, 2173-2181.		2
1231	Evolution of osmosensing OSCA1 Ca ²⁺ channel family coincident with plant transition from water to land. Plant Genome, 2022, 15, e20198.	1.6	5
1232	Pseudo-keriothecal wall texture in Cenomanian Nezzazatina Kaminski 2004: Implications for the higher rank classification of agglutinated foraminifera. Revue De Micropaleontologie, 2022, 75, 100681.	0.8	3
1233	New finds of naked amoebae in the Black Sea (Ukraine). Arxius De Miscellania Zoologica, 0, , 13-25.	0.5	0
1234	A new duplex real-time PCR for simultaneous detection and differentiation of <i>Tetratrichomonas gallinarum</i> and <i>Trichomonas gallinae</i> . Journal of Veterinary Diagnostic Investigation, 2022, , 104063872210980.	0.5	1
1235	Ciliate research: From myth to trendsetting science. Journal of Eukaryotic Microbiology, 2022, 69, .	0.8	3
1237	Gut archaea associated with bacteria colonization and succession during piglet weaning transitions. BMC Veterinary Research, 2022, 18, .	0.7	4

		CITATION REPORT		
#	ARTICLE Influence of aquatic environmental factors on the distribution of naked amoebae in wat	ers of	IF	CITATIONS
1239	Zhytomyr region (Ukraine). Visnyk L'vivs'koho Universytetu Seriia Biolohichna, 2022, , 1		0.0	0
1240	Phylogenomic Analyses of 2,786 Genes in 158 Lineages Support a Root of the Eukaryot between Opisthokonts and All Other Lineages. Genome Biology and Evolution, 2022, 14	ic Tree of Life 4, .	1.1	8
1241	Comentando A Higher Level Classification of All Living Organism y su Correction, de Rug 2015. Ambiociencias, 0, , 102-123.	ggiero, et al.,	0.0	0
1242	Sending the message: specialized RNA export mechanisms in trypanosomes. Trends in F 2022, 38, 854-867.	arasitology,	1.5	6
1243	Giardia duodenalis carries out canonical homologous recombination and single-strand a Research in Microbiology, 2022, 173, 103984.	nnealing.	1.0	0
1244	Identifying groupâ€specific primers for environmental Heterolobosa by highâ€ŧhroughp Microbial Biotechnology, 2022, 15, 2476-2487.	ut sequencing.	2.0	2
1245	Molecular phylogeny of Chonotrichia (Ciliophora, Phyllopharyngea) inferred from SSU rI sequences. European Journal of Protistology, 2022, 86, 125920.	ONA	0.5	1
1246	Seaweed-based fertilizing products. , 2022, , 271-313.			0
1247	Primary production assessment on eco-engineering infrastructures: English Channel cas Conference Series: Materials Science and Engineering, 2022, 1245, 012011.	e study. IOP	0.3	0
1248	Epidemiologia de los coccidios intestinales en personas vulnerables: una revisión sistÃ0 mundial , 2022, 6, 1165-1185.	©mica a nivel		0
1249	Mechanism of action and implication of naphthoquinone as potent anti-trypanosomal d Topics in Medicinal Chemistry, 2022, 22, .	rugs. Current	1.0	0
1250	Largeâ€scale phylogenetic analysis provides insights into the diversification and evolution peritrich ciliates (Protista: Ciliophora). Journal of Eukaryotic Microbiology, 2023, 70, .	on of sessilid	0.8	2
1251	Separate To Operate: the Centriole-Free Inner Core of the Centrosome Regulates the As Intranuclear Spindle in <i>Toxoplasma gondii</i> . MBio, 2022, 13, .	sembly of the	1.8	12
1252	Cryptic and ubiquitous aplastidic cryptophytes are key freshwater flagellated bacterivor Journal, 2023, 17, 84-94.	es. ISME	4.4	3
1253	Sex in protists: A new perspective on the reproduction mechanisms of trypanosomatids Molecular Biology, 2022, 45, .	. Genetics and	0.6	0
1254	Free-Living Amoebas in Extreme Environments: The True Survival in our Planet. BioMed F International, 2022, 2022, 1-10.	Research	0.9	5
1255	Differential Affinity Chromatography Coupled to Mass Spectrometry: A Suitable Tool to Common Binding Proteins of a Broad-Range Antimicrobial Peptide Derived from Leucinc Biomedicines, 2022, 10, 2675.		1.4	8
1256	Pesticide soil microbial toxicity: setting the scene for a new pesticide risk assessment fo microorganisms (IUPAC Technical Report). Pure and Applied Chemistry, 2022, 94, 1161	r soil -1194.	0.9	11

щ		15	CITATION
#	ARTICLE	IF	CITATIONS
1258	Hanging in the air: tree moss diatoms from Indo-Burma biodiversity hot spot of India. Aerobiologia, 2022, 38, 557-566.	0.7	2
1259	The Genome of the Mitochondrion-Related Organelle in Cepedea longa, a Large Endosymbiotic Opalinid Inhabiting the Recta of Frogs. International Journal of Molecular Sciences, 2022, 23, 13472.	1.8	0
1260	In vitro cultivation methods for coccidian parasite research. International Journal for Parasitology, 2023, 53, 477-489.	1.3	5
1261	Protist Diversity Responses to Experimental N Deposition in Biological Crusts of a Semiarid Mediterranean Ecosystem. Protist, 2023, 174, 125929.	0.6	1
1262	Classification of Fungi. , 2023, , 1253-1255.e1.		0
1263	Advances in molecular interactions on the Rhizoctonia solani-sugar beet pathosystem. Fungal Biology Reviews, 2023, 44, 100297.	1.9	2
1264	Proteomic characterization of Toxoplasma gondii ME49 derived strains resistant to the artemisinin derivatives artemiside and artemisone implies potential mode of action independent of ROS formation. International Journal for Parasitology: Drugs and Drug Resistance, 2023, 21, 1-12.	1.4	4
1265	The Transcriptome and Proteome Networks of Malignant Tumours Reveal Atavistic Attractors of Polyploidy-Related Asexual Reproduction. International Journal of Molecular Sciences, 2022, 23, 14930.	1.8	6
1266	Ultrastructure of Diophrys appendiculata and new systematic consideration of the euplotid family Uronychiidae (Protista, Ciliophora). Marine Life Science and Technology, 2022, 4, 551-568.	1.8	7
1267	Shedding light on the origin of Acanthocystidae: Ricksol blepharistes gen. n., sp. n. (Ricksolidae fam.) Tj ETQq1 1 Ozanamia gen. n. (Ozanamiidae fam. n.), and $\hat{a} \in \mathfrak{C}$ Heterophrys-like organisms $\hat{a} \in \mathfrak{C}$ Organisms Diversity and Evolution. O	0.784314 0.7	rgBT /Ove 1
1268	Volatiles of the Apicomplexan Alga <i>Chromera velia</i> and Associated Bacteria. ChemBioChem, 2023, 24, .	1.3	0
1269	Parietal composition of <i>Lichtheimia corymbifera</i> : differences between spore and germ tube stages and host-pathogen interactions. Medical Mycology, 0, , .	0.3	Ο
1270	Essential Oils and Terpenic Compounds as Potential Hits for Drugs against Amitochondriate Protists. Tropical Medicine and Infectious Disease, 2023, 8, 37.	0.9	3
1271	First putative occurrence in the fossil record of choanoflagellates, the sister group of Metazoa. Scientific Reports, 2023, 13, .	1.6	Ο
1273	Infection of the gastrointestinal tract: Giardiasis and amoebiasis. , 2023, , 365-373.		2
1274	Redox partner interactions in the ATG8 lipidation system in microalgae. Free Radical Biology and Medicine, 2023, 203, 58-68.	1.3	2
1276	A three-gene phylogeny supports taxonomic rearrangements in the family Didymiaceae (Myxomycetes). Mycological Progress, 2023, 22, .	0.5	8
1277	Position of Algae on the Tree of Life. Doklady Biological Sciences, 2022, 507, 312-326.	0.2	1

#	Article	IF	Citations
1278	Standard ecological and molecular research methods and techniques for Labyrinthula spp Frontiers in Marine Science, 0, 10, .	1.2	0
1279	Protists: the hidden ecosystem players in a wetland rice field soil. Biology and Fertility of Soils, 0, , .	2.3	1
1280	Taxonomic composition, community structure and molecular novelty of microeukaryotes in a temperate oligomesotrophic lake as revealed by metabarcoding. Scientific Reports, 2023, 13, .	1.6	1
1281	Diversity of algae and their biotechnological potential. Advances in Microbial Physiology, 2023, , 301-321.	1.0	0
1282	The Tubulin Superfamily in Apicomplexan Parasites. Microorganisms, 2023, 11, 706.	1.6	4
1283	Exposure of Culex quinquefasciatus to the oomycete Pythium insidiosum: A protocol for inÂvitro studies. Fungal Biology, 2023, 127, 969-974.	1.1	0
1285	Life cycle and plasmodial axenic culture of <i>Physarum galbeum</i> . Journal of Basic Microbiology, 0,	1.8	0
1286	Identification, Characterization and Use of Microorganisms. , 2023, , 74-119.		0
1287	Mitochondrial Dynamics during Development. , 2023, 2, 19-44.		4
1288	Morphologic and molecular characterization of Apertospathula pilata n. sp., a novel freshwater spathidiid (Ciliophora, Litostomatea) from Idaho, USA. European Journal of Protistology, 2023, 89, 125990.	0.5	1
1298	NADPH Oxidase-Dependent Processes in the Social Amoeba Dictyostelium discoideum. , 2023, , 399-412.		0
1312	The Protistan Origins of Animals and Fungi. , 2023, , 3-38.		0
1314	Zooplankton Communities: Diversity in Time and Space. , 2024, , 539-585.		0
1324	Metagenomics and metatranscriptomics as potential driving forces for the exploration of diversity and functions of micro-eukaryotes in soil. 3 Biotech, 2023, 13, .	1.1	0
1328	Soil fauna: occurrence, biodiversity, and roles in ecosystem function. , 2024, , 131-159.		0
1329	Molecular diagnosis of Toxoplasma gondii. , 2024, , 3049-3060.		0
1336	Algae from Secondary Endosymbiosis. , 2024, , 219-383.		0
1337	Cyanobacteria/Blue-Green Algae. , 2024, , 25-99.		0

ARTICLE

IF CITATIONS