Lei Cai

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/6286794/publications.pdf

Version: 2024-02-01

166	12,835	47006 47	27406 106
papers	citations	h-index	g-index

178	178	178	10456
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Nuclear ribosomal internal transcribed spacer (ITS) region as a universal DNA barcode marker for Fungi. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 6241-6246.	7.1	4,012
2	The Faces of Fungi database: fungal names linked with morphology, phylogeny and human impacts. Fungal Diversity, 2015, 74, 3-18.	12.3	471
3	The Amsterdam Declaration on Fungal Nomenclature. IMA Fungus, 2011, 2, 105-111.	3.8	320
4	Genera of phytopathogenic fungi: GOPHY 1. Studies in Mycology, 2017, 86, 99-216.	7.2	276
5	Resolving the Phoma enigma. Studies in Mycology, 2015, 82, 137-217.	7.2	273
6	One stop shop: backbones trees for important phytopathogenic genera: I (2014). Fungal Diversity, 2014, 67, 21-125.	12.3	241
7	Colletotrichum gloeosporioides is not a common pathogen on tropical fruits. Fungal Diversity, 2010, 44, 33-43.	12.3	225
8	Notes on currently accepted species of Colletotrichum. Mycosphere, 2016, 7, 1192-1260.	6.1	220
9	A phylogenetic and taxonomic re-evaluation of the Bipolaris - Cochliobolus - Curvularia Complex. Fungal Diversity, 2012, 56, 131-144.	12.3	216
10	Naming and outline of Dothideomycetesâ€“2014 including proposals for the protection or suppression of generic names. Fungal Diversity, 2014, 69, 1-55.	12.3	216
11	A multi-locus backbone tree for Pestalotiopsis, with a polyphasic characterization of 14 new species. Fungal Diversity, 2012, 56, 95-129.	12.3	211
12	Fungal diversity notes 929â€“1035: taxonomic and phylogenetic contributions on genera and species of fungi. Fungal Diversity, 2019, 95, 1-273.	12.3	203
13	Didymellaceae revisited. Studies in Mycology, 2017, 87, 105-159.	7.2	172
14	Unravelling <I>Colletotrichum</I> species associated with <I>Camellia</I>: employing ApMat and GS loci to resolve species in the <I>C. gloeosporioides</I> complex. Persoonia: Molecular Phylogeny and Evolution of Fungi, 2015, 35, 63-86.	4.4	166
15	Disease-induced changes in plant microbiome assembly and functional adaptation. Microbiome, 2021, 9, 187.	11.1	157
16	Phylogenetic investigations of Sordariaceae based on multiple gene sequences and morphology. Mycological Research, 2006, 110, 137-150.	2.5	152
17	Application of the Apn2/MAT locus to improve the systematics of the Colletotrichum gloeosporioides complex: an example from coffee (Coffea spp.) hosts. Mycologia, 2012, 104, 396-409.	1.9	152
18	The evolution of species concepts and species recognition criteria in plant pathogenic fungi. Fungal Diversity, 2011, 50, 121-133.	12.3	148

#	Article	IF	Citations
19	Fungal diversity notes 1036â€“1150: taxonomic and phylogenetic contributions on genera and species of fungal taxa. Fungal Diversity, 2019, 96, 1-242.	12.3	148
20	Colletotrichum species causing anthracnose disease of chili in China. Persoonia: Molecular Phylogeny and Evolution of Fungi, 2017, 38, 20-37.	4.4	144
21	Cochliobolus: an overview and current status of species. Fungal Diversity, 2011, 51, 3-42.	12.3	139
22	Phylogenetic reassessment of Nigrospora: ubiquitous endophytes, plant and human pathogens. Persoonia: Molecular Phylogeny and Evolution of Fungi, 2017, 39, 118-142.	4.4	126
23	Species boundaries in plant pathogenic fungi: a Colletotrichum case study. BMC Evolutionary Biology, 2016, 16, 81.	3.2	122
24	Colletotrichum species associated with cultivated citrus in China. Fungal Diversity, 2013, 61, 61-74.	12.3	120
25	Phylogenetics and evolution of nematode-trapping fungi (Orbiliales) estimated from nuclear and protein coding genes. Mycologia, 2005, 97, 1034-1046.	1.9	105
26	Culturable mycobiota from Karst caves in China, with descriptions of 20 new species. Persoonia: Molecular Phylogeny and Evolution of Fungi, 2017, 39, 1-31.	4.4	100
27	Diaporthe is paraphyletic. IMA Fungus, 2017, 8, 153-187.	3.8	100
28	Sporocadaceae, a family of coelomycetous fungi with appendage-bearing conidia. Studies in Mycology, 2019, 92, 287-415.	7.2	94
29	Colletotrichum species from Jasmine (Jasminum sambac). Fungal Diversity, 2011, 46, 171-182.	12.3	90
30	ColletotrichumSpecies onOrchidaceaein Southwest China. Cryptogamie, Mycologie, 2011, 32, 229-253.	1.0	88
31	Diversity, Distribution and Co-occurrence Patterns of Bacterial Communities in a Karst Cave System. Frontiers in Microbiology, 2019, 10, 1726.	3.5	80
32	The <I>Colletotrichum gigasporum</I> species complex. Persoonia: Molecular Phylogeny and Evolution of Fungi, 2014, 33, 83-97.	4.4	79
33	Endophytic Colletotrichum species from Bletilla ochracea (Orchidaceae), with descriptions of seven new speices. Fungal Diversity, 2013, 61, 139-164.	12.3	78
34	How to publish a new fungal species, or name, version 3.0. IMA Fungus, 2021, 12, 11.	3.8	76
35	Unravelling Diaporthe species associated with Camellia. Systematics and Biodiversity, 2016, 14, 102-117.	1.2	73
36	Hostâ€jump drives rapid and recent ecological speciation of the emergent fungal pathogen Colletotrichum kahawae. Molecular Ecology, 2012, 21, 2655-2670.	3.9	72

#	Article	IF	Citations
37	Variation between freshwater and terrestrial fungal communities on decaying bamboo culms. Antonie Van Leeuwenhoek, 2006, 89, 293-301.	1.7	70
38	Two new Kirschsteiniothelia species with Dendryphiopsis anamorphs cluster in Kirschsteiniotheliaceae fam. nov.. Mycologia, 2012, 104, 698-714.	1.9	69
39	Species of the Colletotrichum gloeosporioides complex associated with anthracnose diseases of Proteaceae. Fungal Diversity, 2013, 61, 89-105.	12.3	69
40	Culturable plant pathogenic fungi associated with sugarcane in southern China. Fungal Diversity, 2019, 99, 1-104.	12.3	62
41	Endophytic Colletotrichum from tropical grasses with a new species C. endophytica. Fungal Diversity, 2013, 61, 107-115.	12.3	61
42	An Optimized Protocol of Single Spore Isolation for Fungi. Cryptogamie, Mycologie, 2013, 34, 349-356.	1.0	58
43	Colletotrichum species on grape in Guizhou and Yunnan provinces, China. Mycoscience, 2013, 54, 29-41.	0.8	58
44	Genetic Diversity and Population Structure of Rice Pathogen Ustilaginoidea virens in China. PLoS ONE, 2013, 8, e76879.	2.5	58
45	Pestalotiopsis and allied genera from Camellia, with description of 11 new species from China. Scientific Reports, 2017, 7, 866.	3.3	54
46	Colletotrichum: species, ecology and interactions. IMA Fungus, 2010, 1, 161-165.	3.8	53
47	Culturable mycobiota from Karst caves in China II, with descriptions of 33 new species. Fungal Diversity, 2021, 106, 29-136.	12.3	53
48	The numbers of fungi: contributions from traditional taxonomic studies and challenges of metabarcoding. Fungal Diversity, 2022, 114, 327-386.	12.3	53
49	Morphology: still essential in a molecular world. Mycotaxon, 2011, 114, 439-451.	0.3	52
50	Epitypification of Colletotrichum musae, the causative agent of banana anthracnose. Mycoscience, 2011, 52, 376-382.	0.8	50
51	Eight new Arthrinium species from China. MycoKeys, 2018, 34, 1-24.	1.9	50
52	Psychrophilic fungi from the world's roof. Persoonia: Molecular Phylogeny and Evolution of Fungi, 2015, 34, 100-112.	4.4	49
53	Morphological and molecular characterisation of a new anamorphic genus <I>Cheirosporium</I>, from freshwater in China. Persoonia: Molecular Phylogeny and Evolution of Fungi, 2008, 20, 53-58.	4.4	46
54	Novel Species ofColletotrichumRevealed by Morphology and Molecular Analysis. Cryptogamie, Mycologie, 2012, 33, 347-362.	1.0	46

#	Article	IF	Citations
55	A High-Level Fungal Diversity in the Intertidal Sediment of Chinese Seas Presents the Spatial Variation of Community Composition. Frontiers in Microbiology, 2016, 7, 2098.	3.5	45
56	Phylogeny of Chaetothyriaceae in northern Thailand including three new species. Mycologia, 2012, 104, 382-395.	1.9	44
57	A polyphasic approach to characterise two novel species of Phoma (Didymellaceae) from China . Phytotaxa, 2015, 197, 267-281.	0.3	44
58	Fungal communities in sediments of subtropical Chinese seas as estimated by DNA metabarcoding. Scientific Reports, 2016, 6, 26528.	3.3	43
59	Diaporthe species occurring on Lithocarpus glabra in China, with descriptions of five new species. Fungal Biology, 2015, 119, 295-309.	2.5	42
60	Molecular Systematics of Zopfiella and allied genera: evidence from multi-gene sequence analyses. Mycological Research, 2006, 110, 359-368.	2.5	40
61	Circumscription of the anthracnose pathogens Colletotrichum lindemuthianum and C. nigrum. Mycologia, 2013, 105, 844-860.	1.9	40
62	Fungal Biodiversity Profiles 1â€“10. Cryptogamie, Mycologie, 2015, 36, 121-166.	1.0	40
63	Ten reasons why a sequence-based nomenclature is not useful for fungi anytime soon. IMA Fungus, 2018, 9, 177-183.	3.8	40
64	Three new species of Phomopsis in Gutianshan Nature Reserve in China. Mycological Progress, 2014, 13, 111-121.	1.4	39
65	Species Diversity With Comprehensive Annotations of Wood-Inhabiting Poroid and Corticioid Fungi in Uzbekistan. Frontiers in Microbiology, 2020, 11, 598321.	3.5	39
66	Changes in Bacterial and Fungal Microbiomes Associated with Tomatoes of Healthy and Infected by Fusarium oxysporum f. sp. lycopersici. Microbial Ecology, 2021, 81, 1004-1017.	2.8	39
67	Conlarium duplumascospora gen. et. sp. nov. and Jobellisia guangdongensis sp. nov. from freshwater habitats in China. Mycologia, 2012, 104, 1178-1186.	1.9	38
68	The need to carry out re-inventory of plant pathogenic fungi. Tropical Plant Pathology, 2011, 36, 205-213.	1.5	37
69	3-Anhydro-6-hydroxy-ophiobolin A, a new sesterterpene inhibiting the growth of methicillin-resistant Staphylococcus aureus and inducing the cell death by apoptosis on K562, from the phytopathogenic fungus Bipolaris oryzae. Bioorganic and Medicinal Chemistry Letters, 2013, 23, 3547-3550.	2.2	37
70	Highlighting patterns of fungal diversity and composition shaped by ocean currents using the East China Sea as a model. Molecular Ecology, 2018, 27, 564-576.	3.9	37
71	Introgression and gene family contraction drive the evolution of lifestyle and host shifts of hypocrealean fungi. Mycology, 2018, 9, 176-188.	4.4	35
72	Three new ascomycetes from freshwater in China. Mycologia, 2012, 104, 1478-1489.	1.9	33

#	Article	IF	Citations
73	Utility of Thermostable Xylanases of Mycothermus thermophilus in Generating Prebiotic Xylooligosaccharides. Journal of Agricultural and Food Chemistry, 2017, 65, 1139-1145.	5.2	32
74	Microbiota in the Rhizosphere and Seed of Rice From China, With Reference to Their Transmission and Biogeography. Frontiers in Microbiology, 2020, 11, 995.	3.5	32
75	Oligotrophic fungi from a carbonate cave, with three new species ofCephalotrichum. Mycology, 2017, 8, 164-177.	4.4	31
76	Fungal Community Composition and Potential Depth-Related Driving Factors Impacting Distribution Pattern and Trophic Modes from Epi- to Abyssopelagic Zones of the Western Pacific Ocean. Microbial Ecology, 2019, 78, 820-831.	2.8	31
77	Origin of Cave Fungi. Frontiers in Microbiology, 2018, 9, 1407.	3.5	30
78	Unveiling the Hidden Diversity of Rock-Inhabiting Fungi: Chaetothyriales from China. Journal of Fungi (Basel, Switzerland), 2020, 6, 187.	3.5	30
79	Citizen science project reveals high diversity in Didymellaceae (Pleosporales, Dothideomycetes). MycoKeys, 2020, 65, 49-99.	1.9	29
80	Culture collections, the new herbaria for fungal pathogens. Fungal Diversity, 2010, 45, 21-32.	12.3	28
81	Biology of Colletotrichum horii, the causal agent of persimmon anthracnose. Mycology, 2010, 1, 242-253.	4.4	27
82	A Patatin-Like Protein Associated with the Polyhydroxyalkanoate (PHA) Granules of Haloferax mediterranei Acts as an Efficient Depolymerase in the Degradation of Native PHA. Applied and Environmental Microbiology, 2015, 81, 3029-3038.	3.1	27
83	Phylogeny of new marine Dothideomycetes and Sordariomycetes from mangroves and deep-sea sediments. Botanica Marina, 2020, 63, 155-181.	1.2	27
84	Fungal networks and orchid distribution: new insights from above- and below-ground analyses of fungal communities. IMA Fungus, 2018, 9, 1-11.	3.8	26
85	Polyphasic characterisation of three new Phyllosticta spp.. Persoonia: Molecular Phylogeny and Evolution of Fungi, 2012, 28, 76-84.	4.4	25
86	Cellular and organellar membrane-associated proteins in haloarchaea: Perspectives on the physiological significance and biotechnological applications. Science China Life Sciences, 2012, 55, 404-414.	4.9	25
87	Four new filamentous fungal species from newly-collected and hivestored bee pollen. Mycosphere, 2018, 9, 1089-1116.	6.1	25
88	Biodiversity of aquatic fungi in China. Mycology, 2013, 4, 125-168.	4.4	24
89	Polyphasic characterisation of Chaetomium species from soil and compost revealed high number of undescribed species. Fungal Biology, 2017, 121, 21-43.	2.5	24
90	Distribution of mycotoxin-producing fungi across major rice production areas of China. Food Control, 2022, 134, 108572.	5.5	24

#	Article	IF	Citations
91	Phylogenetic relationships of Chalara and allied species inferred from ribosomal DNA sequences. Mycological Progress, 2009, 8, 133-143.	1.4	23
92	Fungal diversity on submerged wood in a tropical stream and an artificial lake. Biodiversity and Conservation, 2010, 19, 3799-3808.	2.6	23
93	Inferring phylogeny and speciation of Gymnosporangium species and their coevolution with host plants. Scientific Reports, 2016, 6, 29339.	3.3	23
94	A new species of <I>Colletotrichum</I> from <I>Cordyline</I> <I>fruticosa</I> and <I>Eugenia javanica</I> causing anthracnose disease. Mycotaxon, 2011, 114, 247-257.	0.3	22
95	Typification and phylogenetic study of Phyllosticta ampelicida and P. vaccinii. Mycologia, 2013, 105, 1030-1042.	1.9	22
96	Aquapeziza: a new genus from freshwater and its morphological and phylogenetic relationships to Pezizaceae. Mycologia, 2012, 104, 540-546.	1.9	21
97	Substrate and spatial variables are major determinants of fungal community in karst caves in Southwest China. Journal of Biogeography, 2019, 46, 1504-1518.	3.0	21
98	Morphological and Molecular Characterization of a Novel Species ofSimplicilliumfrom China. Cryptogamie, Mycologie, 2012, 33, 137-144.	1.0	20
99	Molecular phylogeny of Neodevriesia, with two new species and several new combinations. Mycologia, 2017, 109, 965-974.	1.9	20
100	Anamorphic fungi from freshwater habitats in China: Dictyosporium tetrasporum and Exserticlava yunnanensis spp. nov., and two new records for Pseudofuscophialis lignicola and Pseudobotrytis terrestris. Mycoscience, 2007, 48, 290-296.	0.8	19
101	Phylogenetic assessment of Chaetomium indicum and allied species, with the introduction of three new species and epitypification of C. funicola and C. indicum. Mycological Progress, 2014, 13, 719-732.	1.4	19
102	Six new soilâ€“inhabiting Cladosporium species from plateaus in China. Mycologia, 2017, 109, 244-260.	1.9	19
103	Temperate Pine Barrens and Tropical Rain Forests Are Both Rich in Undescribed Fungi. PLoS ONE, 2014, 9, e103753.	2.5	18
104	A phylogenetic assessment and taxonomic revision of the thermotolerant hyphomycete generaAcrophialophoraandTaifanglania. Mycologia, 2015, 107, 768-779.	1.9	18
105	Molecular phylogeny ofAscotricha, including two new marine algae-associated species. Mycologia, 2015, 107, 490-504.	1.9	17
106	PenicilliumsectionLanataâ€divaricatafrom acidic soil. Cladistics, 2019, 35, 514-549.	3.3	17
107	A new leaf blight disease of turfgrasses caused by Microdochium poae, sp. nov.. Mycologia, 2019, 111, 265-273.	1.9	17
108	Cryptic fungal species unmasked. Microbiology Australia, 2012, 33, 36.	0.4	17

#	Article	IF	Citations
109	Acrodictys liputii sp. nov. and Digitodesmium bambusicola sp. nov. from bamboo submerged in the Liput River in the Philippines. Nova Hedwigia, 2002, 75, 525-532.	0.4	16
110	Cryptic diversity in Tranzscheliella spp. (Ustilaginales) is driven by host switches. Scientific Reports, 2017, 7, 43549.	3.3	16
111	Species-specific real-time PCR detection ofColletotrichum kahawae. Journal of Applied Microbiology, 2013, 114, 828-835.	3.1	15
112	Cochlioquinone Derivatives with Apoptosisâ€Inducing Effects on HCT116 Colon Cancer Cells from the Phytopathogenic Fungus Bipolaris luttrellii L439. Chemistry and Biodiversity, 2014, 11, 1892-1899.	2.1	15
113	3-Anhydro-6-hydroxy-ophiobolin A_3 a fungal sesterterpene from Bipolaris oryzae induced autophagy and promoted the degradation of Î±-synuclein in PC12 cells. Bioorganic and Medicinal Chemistry Letters, 2015, 25, 1464-1470.	2.2	15
114	Phylogenetic assessment and taxonomic revision of Mariannaea. Mycological Progress, 2017, 16, 271-283.	1.4	15
115	Morphological and molecular characterization of Mariannaea aquaticola sp. nov. collected from freshwater habitats. Mycological Progress, 2010, 9, 337-343.	1.4	14
116	New species and notes of Colletotrichum on daylilies (Hemerocallis spp.). Tropical Plant Pathology, 2012, 37, 165-174.	1.5	14
117	Analysis of the Transcriptional Regulator GlpR, Promoter Elements, and Posttranscriptional Processing Involved in Fructose-Induced Activation of the Phosphoenolpyruvate-Dependent Sugar Phosphotransferase System in Haloferax mediterranei. Applied and Environmental Microbiology, 2014, 80, 1430-1440.	3.1	14
118	Polyphasic characterization of four new plant pathogenic Phyllosticta species from China, Japan, and the United States. Fungal Biology, 2015, 119, 433-446.	2.5	14
119	Polyphasic characterization of Plectosphaerella oligotrophica, aÂnew oligotrophic species from China. Mycoscience, 2013, 54, 387-393.	0.8	13
120	Synopsis ofPhyllostictain China. Mycology, 2015, 6, 50-75.	4.4	13
121	Fungal diversity driven by bark features affects phorophyte preference in epiphytic orchids from southern China. Scientific Reports, 2021, 11, 11287.	3.3	13
122	Temporal and spatial variation of microbial communities in stored rice grains from two major depots in China. Food Research International, 2022, 152, 110876.	6.2	13
123	New species of Clohiesia and Paraniesslia collected from freshwater habitats in China. Mycoscience, 2007, 48, 182-186.	0.8	12
124	Coicenals Aâ€“D, Four New Diterpenoids with New Chemical Skeletons from the Plant Pathogenic Fungus Bipolaris coicis. Organic Letters, 2013, 15, 3982-3985.	4.6	12
125	Polyphasic characterisation of three novel species of Paraboeremia. Mycological Progress, 2017, 16, 285-295.	1.4	12
126	Bacteria and Metabolic Potential in Karst Caves Revealed by Intensive Bacterial Cultivation and Genome Assembly. Applied and Environmental Microbiology, 2021, 87, .	3.1	12

#	Article	IF	Citations
127	Contribution to rust flora in China I, tremendous diversity from natural reserves and parks. Fungal Diversity, 2021, 110, 1-58.	12.3	12
128	Induction of sporulation in plant pathogenic fungi. Mycology, 2012, 3, 195-200.	4.4	12
129	Two new freshwater species of Annulatascaceae from China. Mycotaxon, 2012, 120, 81-88.	0.3	11
130	Three new species of Tilletia on Eriachne from north-western Australia. Mycoscience, 2014, 55, 361-366.	0.8	11
131	Fungal diversity and specificity in Cephalanthera damasonium and C. longifolia (Orchidaceae) mycorrhizas. Journal of Systematics and Evolution, 2017, 55, 158-169.	3.1	11
132	Resolving the Melampsora epitea complex. Mycologia, 2017, 109, 391-407.	1.9	11
133	Seasonal dynamics of mycoplankton in the Yellow Sea reflect the combined effect of riverine inputs and hydrographic conditions. Molecular Ecology, 2021, 30, 3624-3637.	3.9	11
134	Linocarpon bambusicola sp. nov. and Dictyochaeta curvispora sp. nov. from bamboo submerged in freshwater. Nova Hedwigia, 2004, 78, 439-445.	0.4	10
135	RupestriomycesandSpissiomyces, two new genera of rock-inhabiting fungi from China. Mycologia, 2015, 107, 831-844.	1.9	10
136	Halophilic and thermotolerantGymnoascusspecies from several special environments, China. Mycologia, 2016, 108, 179-191.	1.9	10
137	Nanopore sequencing of full <scp>rRNA</scp> operon improves resolution in mycobiome analysis and reveals high diversity in both human gut and environments. Molecular Ecology, 2023, 32, 6330-6344.	3.9	10
138	Rasamsonia composticola, a new thermophilic species isolated from compost in Yunnan, China. Mycological Progress, 2013, 12, 213-221.	1.4	9
139	Myrothecium-like new species from turfgrasses and associated rhizosphere. MycoKeys, 2019, 51, 29-53.	1.9	9
140	Diversity of Pelagic and Benthic Bacterial Assemblages in the Western Pacific Ocean. Frontiers in Microbiology, 2020, 11, 1730.	3.5	9
141	Morphological and phylogenetic characterisation of two new species of Phyllosticta from China. Mycological Progress, 2013, 12, 547-556.	1.4	8
142	Quasipucciniastrum agrimoniae, gen. et sp. nov. on Agrimonia (Rosaceae) from China. Mycology, 2019, 10, 141-150.	4.4	8
143	Establishment of a Genetic Transformation System in Guanophilic Fungus Amphichorda guana. Journal of Fungi (Basel, Switzerland), 2021, 7, 138.	3.5	8
144	Diversity of Moesziomyces (Ustilaginales, Ustilaginomycotina) on Echinochloa and Leersia (Poaceae). MycoKeys, 2019, 52, 1-16.	1.9	8

#	Article	IF	Citations
145	Phylogenetic Relationships, Speciation, and Origin of Armillaria in the Northern Hemisphere: A Lesson Based on rRNA and Elongation Factor 1-Alpha. Journal of Fungi (Basel, Switzerland), 2021, 7, 1088.	3.5	8
146	A Novel Species ofGliocladiopsisfrom Freshwater Habitat in China. Cryptogamie, Mycologie, 2013, 34, 233-241.	1.0	7
147	A new thermophilic species of Myceliophthora from China. Mycological Progress, 2014, 13, 165-170.	1.4	7
148	2H-Pyranone and isocoumarin derivatives isolated from the plant pathogenic fungus Leptosphaena maculans. Journal of Asian Natural Products Research, 2019, 21, 939-946.	1.4	7
149	Impatiens wutaishanensis (Balsaminaceae), a new species from Southeast Yunnan, China. PhytoKeys, 2021, 176, 43-53.	1.0	7
150	Species of Colletotrichum on bamboos from China. Mycologia, 2021, 113, 450-458.	1.9	7
151	Tropical Fungi. Mycology, 2005, , 93-115.	0.5	7
152	Neotypification of Colletotrichum coccodes , the causal agent of potato black dot disease and tomato anthracnose. Mycology, 2011, 2, 248-254.	4.4	6
153	3. The molecular phylogeny of freshwater Sordariomycetes and discomycetes. , 2014, , 47-72.		5
154	Ten new species ofMacalpinomycesonEriachnein northern Australia. Mycologia, 2017, 109, 408-421.	1.9	5
155	Overview of nomenclature novelties of fungi in the world and China (2020). Biodiversity Science, 2021, 29, 1064-1072.	0.6	5
156	Uncovering the mysterious identity of Taisuiâ€"an old Chinese folk legend. Science China Life Sciences, 2020, 63, 1942-1945.	4.9	5
157	Genomics-driven discovery of a new cyclodepsipeptide from the guanophilic fungusAmphichorda guana. Organic and Biomolecular Chemistry, 2021, 19, 1960-1964.	2.8	4
158	Applying early divergent characters in higher rank taxonomy of Melampsorineae (Basidiomycota, Pucciniales). Mycology, 2023, 14, 11-36.	4.4	4
159	Pochonia cordycepisociata, a new species associated with Chinese cordyceps in Tibet, China. Phytotaxa, 2015, 208, 278.	0.3	3
160	Amplisins Aâ€"E, chromone methide polymers with hypoglycemic activity from a new fungicolous fungus Amplistroma fungicola. Organic Chemistry Frontiers, 2020, 7, 2761-2769.	4.5	3
161	Leptosphaerulina species isolated from golf turfgrass in China, with description of L. macrospora, sp. nov.. Mycologia, 2021, 113, 1-12.	1.9	3
162	https://www.fungiofpakistan.com: a continuously updated online database of fungi in Pakistan. Database: the Journal of Biological Databases and Curation, 2021, 2021, .	3.0	2

#	Article	IF	Citations
163	5. Taxonomy of filamentous asexual fungi from freshwater habitats, links to sexual morphs and their phylogeny. , 2014, , 109-132.		1
164	Analysis of macrofungal communities reveals a complex reciprocal influence between Mediterranean montane calcareous grassland and surrounding forest habitats. Journal of Systematics and Evolution, 2021, 59, 278-288.	3.1	1
165	Occurrence and diversity of endophytic fungi in Bletilla ochracea (Orchidaceae) in Guizhou, China. African Journal of Microbiology Research, 2012, 6, .	0.4	1
166	Discovering and dealing with the unknown aspects of Colletotrichum. Mycosphere, 2016, 7, 1074-1075.	6.1	0