

Didymellaceae revisited

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Characterization of fungi causing lesion blight on <i>Papaver dubium</i> in Iran. <i>Antonie Van Leeuwenhoek</i> , 2018, 111, 437-455.	0.7	3
2	<i>Allophoma hayatii</i> sp. nov., an undescribed pathogenic fungus causing dieback of <i>Lantana camara</i> in Iran. <i>Mycological Progress</i> , 2018, 17, 365-379.	0.5	13
3	Coelomycetous <i>Dothideomycetes</i> with emphasis on the families <i>Cucurbitariaceae</i> and <i>Didymellaceae</i> . <i>Studies in Mycology</i> , 2018, 90, 1-69.	4.5	129
4	Tzeananiaceae, a new pleosporalean family associated with <i>Ophiocordyceps macroacicularis</i> fruiting bodies in Taiwan. <i>MycoKeys</i> , 2018, 37, 1-17.	0.8	11
5	The biotechnological potential of <i>Epicoccum</i> spp.: diversity of secondary metabolites. <i>Critical Reviews in Microbiology</i> , 2018, 44, 759-778.	2.7	56
6	Identification and Characterization of Fungal Pathogens Causing Fruit Rot of Deciduous Holly. <i>Plant Disease</i> , 2018, 102, 2430-2445.	0.7	29
7	Fungal Planet description sheets: 716–784. <i>Persoonia: Molecular Phylogeny and Evolution of Fungi</i> , 2018, 40, 239-392.	1.6	142
8	Large-scale generation and analysis of filamentous fungal DNA barcodes boosts coverage for kingdom fungi and reveals thresholds for fungal species and higher taxon delimitation. <i>Studies in Mycology</i> , 2019, 92, 135-154.	4.5	555
9	Whole Genome Sequences of the Tea Leaf Spot Pathogen <i>Didymella segeticola</i> . <i>Phytopathology</i> , 2019, 109, 1676-1678.	1.1	20
10	Genera of phytopathogenic fungi: GOPHY 3. <i>Studies in Mycology</i> , 2019, 94, 1-124.	4.5	104
11	<i>Ectophoma iranica</i> sp. nov. and new hosts and records of <i>Allophoma</i> spp. in Iran. <i>Journal of Phytopathology</i> , 2019, 167, 538-545.	0.5	8
12	Culturable plant pathogenic fungi associated with sugarcane in southern China. <i>Fungal Diversity</i> , 2019, 99, 1-104.	4.7	62
13	Agricultural management and plant selection interactively affect rhizosphere microbial community structure and nitrogen cycling. <i>Microbiome</i> , 2019, 7, 146.	4.9	202
14	Fungal diversity notes 1036–1150: taxonomic and phylogenetic contributions on genera and species of fungal taxa. <i>Fungal Diversity</i> , 2019, 96, 1-242.	4.7	148
15	Host metabolite producing endophytic fungi isolated from <i>Hypericum perforatum</i> . <i>PLoS ONE</i> , 2019, 14, e0217060.	1.1	32
16	<i>Neosascochyta</i> species cause leaf scorch on wheat in Australia. <i>Australasian Plant Disease Notes</i> , 2019, 14, 1.	0.4	13
17	<i>Epicoccum layuense</i> a potential biological control agent of esca-associated fungi in grapevine. <i>PLoS ONE</i> , 2019, 14, e0213273.	1.1	47
18	One stop shop II: taxonomic update with molecular phylogeny for important phytopathogenic genera: 26–50 (2019). <i>Fungal Diversity</i> , 2019, 94, 41-129.	4.7	69

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19	Identification of a Polyketide Synthase Gene Responsible for Ascochitine Biosynthesis in <i>Ascochyta fabae</i> and Its Abrogation in Sister Taxa. <i>MSphere</i> , 2019, 4, .	1.3	6
20	In vitro inferred interactions of selected entomopathogenic fungi from Taiwan and eggs of <i>Meloidogyne graminicola</i> . <i>Mycological Progress</i> , 2020, 19, 97-109.	0.5	9
21	First Report of <i>Epicoccum layuense</i> Causing Brown Leaf Spot on Oat (<i>Avena sativa</i>) in Northwestern China. <i>Plant Disease</i> , 2020, 104, 990-990.	0.7	4
22	Reconsideration of species boundaries and proposed DNA barcodes for <i>Calonectria</i> . <i>Studies in Mycology</i> , 2020, 97, 100106.	4.5	39
23	Taxonomic novelties in Magnolia-associated pleosporalean fungi in the Kunming Botanical Gardens (Yunnan, China). <i>PLoS ONE</i> , 2020, 15, e0235855.	1.1	35
24	Distribution of fungi and their toxic metabolites in melon and sesame seeds marketed in two major producing states in Nigeria. <i>Mycotoxin Research</i> , 2020, 36, 361-369.	1.3	10
25	Evaluating the Microbiome of Hemp. <i>Phytobiomes Journal</i> , 2020, 4, 351-363.	1.4	12
26	Dynamic Changes in the Microbiome of Rice During Shoot and Root Growth Derived From Seeds. <i>Frontiers in Microbiology</i> , 2020, 11, 559728.	1.5	29
27	Five Fungal Pathogens Are Responsible for Bayberry Twig Blight and Fungicides Were Screened for Disease Control. <i>Microorganisms</i> , 2020, 8, 689.	1.6	17
28	The phoma-like dilemma. <i>Studies in Mycology</i> , 2020, 96, 309-396.	4.5	87
29	Fungal diversity notes 1151–1276: taxonomic and phylogenetic contributions on genera and species of fungal taxa. <i>Fungal Diversity</i> , 2020, 100, 5-277.	4.7	156
30	First Report of Leaf Spot Disease Caused by <i>Epicoccum layuense</i> on <i>Camellia sinensis</i> in Chongqing, China. <i>Plant Disease</i> , 2020, 104, 2029.	0.7	11
31	<i>Didymella corylicola</i> sp. nov., a new fungus associated with hazelnut fruit development in Italy. <i>Mycological Progress</i> , 2020, 19, 317-328.	0.5	11
32	Taxonomy and phylogeny of hyaline-spored coelomycetes. <i>Fungal Diversity</i> , 2020, 100, 279-801.	4.7	58
33	Characterization of the fungal community in the canopy air of the invasive plant <i>Ageratina adenophora</i> and its potential to cause plant diseases. <i>PLoS ONE</i> , 2020, 15, e0230822.	1.1	7
34	Microfungi associated with <i>Clematis</i> (Ranunculaceae) with an integrated approach to delimiting species boundaries. <i>Fungal Diversity</i> , 2020, 102, 1-203.	4.7	93
35	Fungal diseases of non-conventional food plants: first report of <i>Stagonosporopsis caricae</i> causing leaf spots on <i>Vasconcellea monoica</i> . <i>Australasian Plant Disease Notes</i> , 2020, 15, 1.	0.4	6
36	Molecular and Morphological Characterization of Two Novel Species Collected from Soil in Korea. <i>Mycobiology</i> , 2020, 48, 9-19.	0.6	6

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37	Taxonomic and phylogenetic contributions to fungi associated with the invasive weed <i>Chromolaena odorata</i> (Siam weed). <i>Fungal Diversity</i> , 2020, 101, 1-175.	4.7	82
38	Fungal Diversity and Mycotoxins in Low Moisture Content Ready-To-Eat Foods in Nigeria. <i>Frontiers in Microbiology</i> , 2020, 11, 615.	1.5	22
39	The Fungal Endophyte <i>Epicoccum dendrobii</i> as a Potential Biocontrol Agent Against <i>Colletotrichum gloeosporioides</i> . <i>Phytopathology</i> , 2021, 111, 293-303.	1.1	30
40	Identification and Characterization of <i>Nothophoma quercina</i> Causing Bud Blight on <i>Photinia</i> – <i>fraseri</i> in China. <i>Plant Disease</i> , 2021, 105, 1356-1364.	0.7	5
41	Change in root-associated fungal communities affects soil enzymatic activities during <i>Pinus massoniana</i> forest development in subtropical China. <i>Forest Ecology and Management</i> , 2021, 482, 118817.	1.4	27
42	Leaf Spot Caused by <i>Epicoccum laticollum</i> on Tobacco in China. <i>Plant Disease</i> , 2021, 105, 501-501.	0.7	7
44	<i>Remotididymella ageratinae</i> sp. nov. and <i>Remotididymella anemophila</i> sp. nov., two novel species isolated from the invasive weed <i>Ageratina adenophora</i> in PR China. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2021, 71, .	0.8	4
45	Additions to Italian Pleosporinae, including <i>Italica heraclei</i> sp. nov.. <i>Biodiversity Data Journal</i> , 2021, 9, e59648.	0.4	1
46	Integrated mRNA and Small RNA Sequencing for Analyzing Leaf Spot Pathogen <i>Didymella segeticola</i> and Its Host, Tea (<i>Camellia sinensis</i>), During Infection. <i>Molecular Plant-Microbe Interactions</i> , 2021, 34, 127-130.	1.4	11
47	Microbial community analysis of soils under different soybean cropping regimes in the Argentinean south-eastern Humid Pampas. <i>FEMS Microbiology Ecology</i> , 2021, 97, .	1.3	12
48	First Report of <i>Epicoccum tobaicum</i> Associated with Leaf Spot on Flowering Cherry in South Korea. <i>Plant Disease</i> , 2021, , .	0.7	4
49	Identification and Characterization of Leaf-Inhabiting Fungi from <i>Castanea</i> Plantations in China. <i>Journal of Fungi</i> (Basel, Switzerland), 2021, 7, 64.	1.5	38
50	Outline of Ascomycota. , 2021, , 246-254.		5
51	<i>Epicoccum</i> species: ubiquitous plant pathogens and effective biological control agents. <i>European Journal of Plant Pathology</i> , 2021, 159, 713-725.	0.8	36
52	Cryptic diversity found in Didymellaceae from Australian native legumes. <i>MycKeys</i> , 2021, 78, 1-20.	0.8	13
53	<i>Allophoma</i> species (Pleosporales: Didymellaceae) associated with <i>Thunbergia grandiflora</i> in Guangxi Province, China. <i>Biodiversity Data Journal</i> , 2021, 9, e63643.	0.4	5
54	Fungi in PAH-contaminated marine sediments: Cultivable diversity and tolerance capacity towards PAH. <i>Marine Pollution Bulletin</i> , 2021, 164, 112082.	2.3	22
55	Investigation of Fungal Strains Composition in Fruit Pollens for Artificial Pollination. <i>Mycobiology</i> , 2021, 49, 249-257.	0.6	2

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56	Whole Genome Sequence and Gene Annotation Resource for <i>Didymella bellidis</i> Associated with Tea Leaf Spot. <i>Plant Disease</i> , 2021, 105, 1168-1170.	0.7	3
57	Revealing of biodiversity and antimicrobial effects of <i>Artemisia asiatica</i> endophytes. <i>Acta Biologica Szegediensis</i> , 2021, 64, 111-119.	0.7	0
58	Isolation and Molecular Characterization of the Romaine Lettuce Phylloplane Mycobiome. <i>Journal of Fungi (Basel, Switzerland)</i> , 2021, 7, 277.	1.5	11
59	New Coelomycetous Fungi from Freshwater in Spain. <i>Journal of Fungi (Basel, Switzerland)</i> , 2021, 7, 368.	1.5	9
60	<i>Sphaeropsis sapinea</i> and Associated Endophytes in Scots Pine: Interactions and Effect on the Host Under Variable Water Content. <i>Frontiers in Forests and Global Change</i> , 2021, 4, .	1.0	19
61	The bacterial and fungal nest microbiomes in populations of the social spider <i>Stegodyphus dumicola</i> . <i>Systematic and Applied Microbiology</i> , 2021, 44, 126222.	1.2	12
62	First Report of <i>Epicoccum nigrum</i> Causing Brown Leaf Spot in Tea in Guizhou Province, China. <i>Plant Disease</i> , 2022, 106, 321.	0.7	11
63	Leaf Blight of <i>Magnolia coco</i> Caused by <i>Nothophoma quercina</i> in China. <i>Plant Disease</i> , 2022, 106, 761.	0.7	3
64	Species concepts of Dothideomycetes: classification, phylogenetic inconsistencies and taxonomic standardization. <i>Fungal Diversity</i> , 2021, 109, 283-319.	4.7	26
65	The Sequence and Integrated Analysis of Competing Endogenous RNAs Originating from Tea Leaves Infected by the Pathogen of Tea Leaf Spot, <i>Didymella segeticola</i> . <i>Plant Disease</i> , 2022, 106, 1286-1290.	0.7	4
66	First Characterisation of the Phoma Species Complex on Maize Leaves in Central Europe. <i>Pathogens</i> , 2021, 10, 1216.	1.2	1
67	First Report of Leaf Spot of <i>Weigela florida</i> Caused by <i>Epicoccum layuense</i> in China. <i>Plant Disease</i> , 2021, , PDIS-07-20-1498.	0.7	2
68	First report of leaf spot on <i>Elaeagnus pungens</i> caused by <i>Epicoccum laticollum</i> in China. <i>Forest Pathology</i> , 2021, 51, e12716.	0.5	8
69	Retrotransposons and multilocus sequence analysis reveals diversity and genetic variability in endophytic fungi-associated with <i>Serjania laruooteana</i> Cambess. <i>Brazilian Journal of Microbiology</i> , 2021, 52, 2179-2192.	0.8	1
70	Uncovering the hidden taxonomic diversity of fungi in Oman. <i>Fungal Diversity</i> , 2021, 106, 229-268.	4.7	11
71	Exploring Fungal Biodiversity of Genus <i>Epicoccum</i> and Their Biotechnological Potential. <i>Fungal Biology</i> , 2021, , 237-276.	0.3	0
72	One stop shop III: taxonomic update with molecular phylogeny for important phytopathogenic genera: 51â€™75 (2019). <i>Fungal Diversity</i> , 2019, 98, 77-160.	4.7	35
73	Species Identification in Plant-Associated Prokaryotes and Fungi Using DNA. <i>Phytobiomes Journal</i> , 2020, 4, 103-114.	1.4	7

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74	First Report of Leaf Blight on <i>Aucuba japonica</i> Caused by <i>Nothophoma quercina</i> in China. <i>Plant Disease</i> , 2020, 104, 2731-2731.	0.7	3
75	<i>Verrucoconiothyrium ambiguum</i> sp. nov., a novel species isolated from sea water, and affiliation of the genus <i>Verrucoconiothyrium</i> to the family <i>Didymellaceae</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2019, 69, 3769-3776.	0.8	7
76	Revealing the hidden diversity of marine fungi in Portugal with the description of two novel species, <i>Neoscochyta fuci</i> sp. nov. and <i>Paraconiothyrium salinum</i> sp. nov.. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020, 70, 5337-5354.	0.8	17
77	Phoma diseases: Epidemiology and control. <i>Plant Pathology</i> , 2020, 69, 1203-1217.	1.2	39
78	Fungal Systematics and Evolution: FUSE 5. <i>Sydowia</i> , 2019, 71, 141-245.	3.7	24
79	Swim bladder mycosis in farmed rainbow trout <i>Oncorhynchus mykiss</i> caused by <i>Phoma herbarum</i> and experimental verification of pathogenicity. <i>Diseases of Aquatic Organisms</i> , 2020, 138, 237-246.	0.5	2
80	Citizen science project reveals high diversity in <i>Didymellaceae</i> (Pleosporales, Dothideomycetes). <i>MycKeys</i> , 2020, 65, 49-99.	0.8	29
81	Diversity and toxigenicity of fungi and description of <i>Fusarium madaense</i> sp. nov. from cereals, legumes and soils in north-central Nigeria. <i>MycKeys</i> , 2020, 67, 95-124.	0.8	20
82	First report of <i>Epicoccum nigrum</i> causing disease in <i>Lotus corniculatus</i> in Argentina. <i>New Disease Reports</i> , 2018, 38, 6-6.	0.4	7
83	Microbial diversity and associated metabolic potential in the supraglacial habitat of a fast-retreating glacier: a case study of Patsio glacier, North-western Himalaya. <i>Environmental Microbiology Reports</i> , 2022, 14, 443-452.	1.0	5
85	First Report of <i>Didymella americana</i> on Lima Bean (<i>Phaseolus lunatus</i>) in Delaware and Maryland. <i>Plant Disease</i> , 2020, 104, 1860-1860.	0.7	4
86	The mycobiome of the oral cavity in healthy dogs and dogs with periodontal disease. <i>American Journal of Veterinary Research</i> , 2022, 83, 42-49.	0.3	7
89	Catnip (<i>Nepeta cataria</i> L.): Recent Advances in , Horticulture and Production. <i>Medicinal and Aromatic Plants of the World</i> , 2020, , 247-284.	0.1	4
90	First Report of Stem Spots of <i>Ephedra intermedia</i> Caused by <i>Neodidymelliopsis longicolla</i> . <i>Plant Disease</i> , 2020, 104, 1253.	0.7	1
91	First Report of <i>Didymella bellidis</i> Causing Tea Leaf Spot in China. <i>Plant Disease</i> , 2020, 104, 1254-1254.	0.7	9
93	Taxonomical Evaluation of Phoma: History of Classification, Current Status and Future Directions. , 2022, , 13-34.		0
94	Overview of Phoma-Like Fungi on Important Legumes (Papilionaceous Plants). , 2022, , 65-89.		1
95	Fungal Planet description sheets: 1284-1382. <i>Persoonia: Molecular Phylogeny and Evolution of Fungi</i> , 2021, 47, 178-374.	1.6	44

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96	Temporal and spatial variation of microbial communities in stored rice grains from two major depots in China. <i>Food Research International</i> , 2022, 152, 110876.	2.9	13
97	Fungal diversity on brewery filling hall surfaces and quality control samples. <i>Yeast</i> , 2022, 39, 141-155.	0.8	7
98	<i>Xenodidymella iranica</i> sp. nov. and new hosts of <i>X. glycyrrhizicola</i> in Iran. <i>Tropical Plant Pathology</i> , 0, 1.	0.8	0
99	Weeds Harbor an Impressive Diversity of Fungi, Which Offers Possibilities for Biocontrol. <i>Applied and Environmental Microbiology</i> , 2022, 88, aem0217721.	1.4	4
100	First Report of Maize Stalk Rot Caused by <i>Epicoccum laticollum</i> on Maize (<i>Zea mays</i>) in China. <i>Plant Disease</i> , 2022, 106, 2255.	0.7	6
101	Endophytic Fungi Isolated from <i>Baccharis linearis</i> and <i>Echinopsis chiloensis</i> with Antifungal Activity against <i>Botrytis cinerea</i> . <i>Journal of Fungi (Basel, Switzerland)</i> , 2022, 8, 197.	1.5	5
102	New species of the family Didymellaceae in Iran. <i>Mycological Progress</i> , 2022, 21, 1.	0.5	4
103	Current Insight into Traditional and Modern Methods in Fungal Diversity Estimates. <i>Journal of Fungi (Basel, Switzerland)</i> , 2022, 8, 226.	1.5	20
104	Multi-Omics Analysis Reveals that the Antimicrobial Kasugamycin Potential Targets Nitrate Reductase in <i>Didymella segeticola</i> to Achieve Control of Tea Leaf Spot. <i>Phytopathology</i> , 2022, 112, 1894-1906.	1.1	5
105	<i>Epicoccum layuense</i> Causing Leaf Spot on <i>Oxalis corymbosa</i> in China. <i>Plant Disease</i> , 2022, 106, 2992.	0.7	3
106	Forecasting the number of species of asexually reproducing fungi (Ascomycota and Basidiomycota). <i>Fungal Diversity</i> , 2022, 114, 463-490.	4.7	12
107	A Culture-Based Study of Micromycetes Isolated from the Urban Nests of Grey Heron (<i>Ardea cinerea</i>) in SW Poland. <i>Animals</i> , 2022, 12, 676.	1.0	3
108	Fungal diversity notes 1387â€“1511: taxonomic and phylogenetic contributions on genera and species of fungal taxa. <i>Fungal Diversity</i> , 2021, 111, 1-335.	4.7	88
109	New Dothideomycetes from Freshwater Habitats in Spain. <i>Journal of Fungi (Basel, Switzerland)</i> , 2021, 7, 1102.	1.5	7
110	A Novel Real Time PCR Method for the Detection and Quantification of <i>Didymella pinodella</i> in Symptomatic and Asymptomatic Plant Hosts. <i>Journal of Fungi (Basel, Switzerland)</i> , 2022, 8, 41.	1.5	7
114	Virulence and Host Range of Fungi Associated With the Invasive Plant <i>Ageratina adenophora</i> . <i>Frontiers in Microbiology</i> , 2022, 13, 857796.	1.5	2
115	Fungal-fungal cocultivation leads to widespread secondary metabolite alteration requiring the partial loss-of-function VeA1 protein. <i>Science Advances</i> , 2022, 8, eabo6094.	4.7	27
116	First Report of <i>Epicoccum nigrum</i> Causing Leaf Spot of <i>Eugenia involucrata</i> in Brazil. <i>Plant Disease</i> , 2023, 107, 230.	0.7	2

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117	<i>Epicoccum</i> spp. Causing Maize Leaf Spot in Heilongjiang Province, China. <i>Plant Disease</i> , 2022, 106, 3050-3060.	0.7	1
118	Genera of phytopathogenic fungi: GOPHY 4. <i>Studies in Mycology</i> , 2022, 101, 417-564.	4.5	36
119	Amplicon Sequencing Reveals Novel Fungal Species Responsible for a Controversial Tea Disease. <i>Journal of Fungi (Basel, Switzerland)</i> , 2022, 8, 782.	1.5	4
120	Functional Annotation, Prediction of <i>Cis</i> Target Gene for the Sequence of mRNAs, and Candidate Long Noncoding RNAs from Tea (<i>Camellia sinensis</i> var. <i>sinensis</i>) Leaves During Infection by the Fungal Pathogen <i>Epicoccum nigrum</i> . <i>PhytoFrontiers</i> , 0, , .	0.8	0
121	Diversity and Pathogenicity of Fungi Associated with Fruit Rot of Winter Jujube in Shandong Province, China. <i>Plant Disease</i> , 2023, 107, 794-801.	0.7	1
122	Integrated Sequencing Data, Annotation, and Targeting Analysis of mRNAs and MicroRNAs from Tea Leaf During Infection by Tea Leaf Spot Pathogen, <i>Epicoccum nigrum</i> . <i>Plant Disease</i> , 2022, 106, 2741-2745.	0.7	2
123	Fungal endophyte <i>Epicoccum nigrum</i> 38L1 inhibits in vitro and in vivo the pathogenic fungus <i>Fusarium graminearum</i> . <i>Biological Control</i> , 2022, 174, 105010.	1.4	9
124	Seed pathology of non-domesticated species of tropical ecosystems. <i>Journal of Seed Science</i> , 0, 44, .	0.7	1
125	Integrated Transcriptome and Proteome Analysis Reveals that the Antimicrobial Griseofulvin Targets <i>Didymella segeticola</i> beta-Tubulin to Control Tea Leaf Spot. <i>Phytopathology</i> , 0, , .	1.1	1
126	Sequence Data, Functional Annotation, and Relationship Analysis Between mRNAs and Long Noncoding RNAs from Tea Leaves During Infection by the Fungal Pathogen <i>Epicoccum sorghinum</i> . <i>Molecular Plant-Microbe Interactions</i> , 2022, 35, 875-879.	1.4	0
127	Survey and Identification of Didymellaceae Causing Stem Canker Disease of Eucalyptus (E.) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 342 T	0.3	0
128	Long-term fungus-plant covariation from multi-site sedimentary ancient DNA metabarcoding. <i>Quaternary Science Reviews</i> , 2022, 295, 107758.	1.4	5
130	Screening for broad-spectrum antimicrobial endophytes from <i>Rosa roxburghii</i> and multi-omic analyses of biosynthetic capacity. <i>Frontiers in Plant Science</i> , 0, 13, .	1.7	5
131	First Report of Brown Leaf Spot Caused by <i>Epicoccum tobaicum</i> on Oat (<i>Avena sativa</i>) in Korea. <i>Plant Disease</i> , 2023, 107, 2255.	0.7	1
132	Plants control the structure of mycorrhizal and pathogenic fungal communities in soil in a 50-year maize monoculture experiment. <i>Plant and Soil</i> , 2023, 484, 133-153.	1.8	5
133	Isolation and identification of <i>Epicoccum nigrum</i> as the causal agent of brown spot disease in <i>Solanum tuberosum</i> in China. <i>Plant Pathology</i> , 2023, 72, 829-838.	1.2	2
134	Revision of the <i>Microsphaeropsis</i> Complex with Addition of Four New <i>Paramicrosphaeropsis</i> L.W.Hou, L.Cai & Crous Species from Zagrosian Forest Trees in Iran. <i>Cryptogamie, Mycologie</i> , 2022, 43, .	0.2	2
135	Genome and Transcriptome Analysis of <i>Ascochyta pisi</i> Provides Insights into the Pathogenesis of <i>Ascochyta</i> Blight of Pea. <i>Microbiology Spectrum</i> , 2023, 11, .	1.2	2

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136	Identification and Characterization of Foliar Fungi Associated with Beach Vitex (<i>Vitex rotundifolia</i> L.). <i>Forests</i> , 2023, 14, 220.	0.9	0
137	Plant-Associated Novel Didymellaceous Taxa in the South China Botanical Garden (Guangzhou, China). <i>Journal of Fungi</i> (Basel, Switzerland), 2023, 9, 182.	1.5	5
138	Changes in soil microbial diversity and community composition across bahiagrass and rhizoma peanut pastures. <i>Biology and Fertility of Soils</i> , 2023, 59, 285-300.	2.3	1
139	First Report of <i>Epicoccum laticollum</i> Causing Leaf Blight on <i>Curcuma kwangsiensis</i> in China. <i>Plant Disease</i> , 2023, 107, 2546.	0.7	1
140	Tapping Culture Collections for Fungal Endophytes: First Genome Assemblies for Three Genera and Five Species in the <i>Ascomycota</i> . <i>Genome Biology and Evolution</i> , 2023, 15, .	1.1	3
141	Two Novel Species of <i>Mesophoma</i> gen. nov. from China. <i>Current Microbiology</i> , 2023, 80, .	1.0	1
142	Characteristics of Leaf Spot Disease Caused by <i>Didymella</i> Species and the Influence of Infection on Tea Quality. <i>Phytopathology</i> , 2023, 113, 516-527.	1.1	8
143	First Report of <i>Epicoccum laticollum</i> Causing Leaf Spot Disease on Banana in China. <i>Plant Disease</i> , 2023, 107, 2553.	0.7	1
150	<i>Phoma</i> spp. an untapped treasure of cytotoxic compounds: current status and perspectives. <i>Applied Microbiology and Biotechnology</i> , 2023, 107, 4991-5001.	1.7	1