

# David S Hibbett

## List of Publications by Year in descending order

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131  
papers

23,221  
citations

11608

70  
h-index

12910

131  
g-index

136  
all docs

136  
docs citations

136  
times ranked

14639  
citing authors

#	ARTICLE	IF	CITATIONS
1	A higher-level phylogenetic classification of the Fungi. <i>Mycological Research</i> , 2007, 111, 509-547.	2.5	1,994
2	Reconstructing the early evolution of Fungi using a six-gene phylogeny. <i>Nature</i> , 2006, 443, 818-822.	13.7	1,625
3	Phylogenetic Species Recognition and Species Concepts in Fungi. <i>Fungal Genetics and Biology</i> , 2000, 31, 21-32.	0.9	1,585
4	The Paleozoic Origin of Enzymatic Lignin Decomposition Reconstructed from 31 Fungal Genomes. <i>Science</i> , 2012, 336, 1715-1719.	6.0	1,424
5	Convergent losses of decay mechanisms and rapid turnover of symbiosis genes in mycorrhizal mutualists. <i>Nature Genetics</i> , 2015, 47, 410-415.	9.4	870
6	Assembling the fungal tree of life: progress, classification, and evolution of subcellular traits. <i>American Journal of Botany</i> , 2004, 91, 1446-1480.	0.8	718
7	Extensive sampling of basidiomycete genomes demonstrates inadequacy of the white-rot/brown-rot paradigm for wood decay fungi. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 9923-9928.	3.3	595
8	Synthesis of phylogeny and taxonomy into a comprehensive tree of life. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 12764-12769.	3.3	584
9	Genome, transcriptome, and secretome analysis of wood decay fungus <i>Postia placenta</i> supports unique mechanisms of lignocellulose conversion. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 1954-1959.	3.3	530
10	The Plant Cell Wallâ€“Decomposing Machinery Underlies the Functional Diversity of Forest Fungi. <i>Science</i> , 2011, 333, 762-765.	6.0	512
11	Major clades of Agaricales: a multilocus phylogenetic overview. <i>Mycologia</i> , 2006, 98, 982-995.	0.8	449
12	Evolutionary instability of ectomycorrhizal symbioses in basidiomycetes. <i>Nature</i> , 2000, 407, 506-508.	13.7	426
13	Genome sequence of the button mushroom <i>Agaricus bisporus</i> reveals mechanisms governing adaptation to a humic-rich ecological niche. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 17501-17506.	3.3	359
14	Contributions of <i>rpb2</i> and <i>tef1</i> to the phylogeny of mushrooms and allies (Basidiomycota, Fungi). <i>Molecular Phylogenetics and Evolution</i> , 2007, 43, 430-451.	1.2	341
15	The phylogenetic distribution of resupinate forms across the major clades of mushroomâ€“forming fungi (Homobasidiomycetes). <i>Systematics and Biodiversity</i> , 2005, 3, 113-157.	0.5	340
16	The Amsterdam Declaration on Fungal Nomenclature. <i>IMA Fungus</i> , 2011, 2, 105-111.	1.7	320
17	Unearthing the roots of ectomycorrhizal symbioses. <i>Nature Reviews Microbiology</i> , 2016, 14, 760-773.	13.6	317
18	Progress in molecular and morphological taxon discovery in Fungi and options for formal classification of environmental sequences. <i>Fungal Biology Reviews</i> , 2011, 25, 38-47.	1.9	296

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19	The Fungi. <i>Current Biology</i> , 2009, 19, R840-R845.	1.8	279
20	Major clades of Agaricales: a multilocus phylogenetic overview. <i>Mycologia</i> , 2006, 98, 982-995.	0.8	268
21	Ectomycorrhizal fungi decompose soil organic matter using oxidative mechanisms adapted from saprotrophic ancestors. <i>New Phytologist</i> , 2016, 209, 1705-1719.	3.5	264
22	Comparative genomics of <i>Ceriporiopsis subvermispora</i> and <i>Phanerochaete chrysosporium</i> provide insight into selective ligninolysis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 5458-5463.	3.3	259
23	Phylogenetic and phylogenomic overview of the Polyporales. <i>Mycologia</i> , 2013, 105, 1350-1373.	0.8	259
24	Large-scale genome sequencing of mycorrhizal fungi provides insights into the early evolution of symbiotic traits. <i>Nature Communications</i> , 2020, 11, 5125.	5.8	258
25	Molecular systematics and biological diversification of Boletales. <i>Mycologia</i> , 2006, 98, 971-981.	0.8	215
26	Comparative Genomics of Early-Diverging Mushroom-Forming Fungi Provides Insights into the Origins of Lignocellulose Decay Capabilities. <i>Molecular Biology and Evolution</i> , 2016, 33, 959-970.	3.5	213
27	A revised family-level classification of the Polyporales (Basidiomycota). <i>Fungal Biology</i> , 2017, 121, 798-824.	1.1	190
28	Contemporaneous radiations of fungi and plants linked to symbiosis. <i>Nature Communications</i> , 2018, 9, 5451.	5.8	189
29	Megaphylogeny resolves global patterns of mushroom evolution. <i>Nature Ecology and Evolution</i> , 2019, 3, 668-678.	3.4	187
30	Out of the Palaeotropics? Historical biogeography and diversification of the cosmopolitan ectomycorrhizal mushroom family Inocybaceae. <i>Journal of Biogeography</i> , 2009, 36, 577-592.	1.4	184
31	Evolution of complex fruiting-body morphologies in homobasidiomycetes. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2002, 269, 1963-1969.	1.2	179
32	A phylogenetic overview of the Agaricomycotina. <i>Mycologia</i> , 2006, 98, 917-925.	0.8	172
33	Fungal functional ecology: bringing a trait-based approach to plant-associated fungi. <i>Biological Reviews</i> , 2020, 95, 409-433.	4.7	171
34	Analysis of Character Correlations Among Wood Decay Mechanisms, Mating Systems, and Substrate Ranges in Homobasidiomycetes. <i>Systematic Biology</i> , 2001, 50, 215-242.	2.7	170
35	Molecular systematics and biological diversification of Boletales. <i>Mycologia</i> , 2006, 98, 971-981.	0.8	167
36	Evolution of helotialean fungi (Leotiomyces, Pezizomycotina): A nuclear rDNA phylogeny. <i>Molecular Phylogenetics and Evolution</i> , 2006, 41, 295-312.	1.2	165

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37	Amylocorticiales ord. nov. and Jaapiales ord. nov.: Early diverging clades of Agaricomycetidae dominated by corticioid forms. <i>Mycologia</i> , 2010, 102, 865-880.	0.8	165
38	Analysis of Character Correlations Among Wood Decay Mechanisms, Mating Systems, and Substrate Ranges in Homobasidiomycetes. <i>Systematic Biology</i> , 2001, 50, 215-242.	2.7	161
39	Sequence-based classification and identification of Fungi. <i>Mycologia</i> , 2016, 108, 1049-1068.	0.8	154
40	Fungal systematics: is a new age of enlightenment at hand?. <i>Nature Reviews Microbiology</i> , 2013, 11, 129-133.	13.6	153
41	The relative ages of ectomycorrhizal mushrooms and their plant hosts estimated using Bayesian relaxed molecular clock analyses. <i>BMC Biology</i> , 2009, 7, 13.	1.7	152
42	An overview of the higher level classification of Pucciniomycotina based on combined analyses of nuclear large and small subunit rDNA sequences. <i>Mycologia</i> , 2006, 98, 896-905.	0.8	143
43	Phylogenetic overview of the Boletineae. <i>Fungal Biology</i> , 2013, 117, 479-511.	1.1	143
44	Evolution of novel wood decay mechanisms in Agaricales revealed by the genome sequences of <i>Fistulina hepatica</i> and <i>Cylindrobasidium torrendii</i> . <i>Fungal Genetics and Biology</i> , 2015, 76, 78-92.	0.9	141
45	Higher-Level Phylogenetic Relationships of Homobasidiomycetes (Mushroom-Forming Fungi) Inferred from Four rDNA Regions. <i>Molecular Phylogenetics and Evolution</i> , 2002, 22, 76-90.	1.2	140
46	The search for the fungal tree of life. <i>Trends in Microbiology</i> , 2009, 17, 488-497.	3.5	139
47	Lignin-degrading peroxidases in Polyporales: an evolutionary survey based on 10 sequenced genomes. <i>Mycologia</i> , 2013, 105, 1428-1444.	0.8	134
48	Latent homology and convergent regulatory evolution underlies the repeated emergence of yeasts. <i>Nature Communications</i> , 2014, 5, 4471.	5.8	133
49	Progress toward a phylogenetic classification of the Polyporaceae through parsimony analysis of mitochondrial ribosomal DNA sequences. <i>Canadian Journal of Botany</i> , 1995, 73, 853-861.	1.2	132
50	Fossil mushrooms from Miocene and Cretaceous ambers and the evolution of Homobasidiomycetes. <i>American Journal of Botany</i> , 1997, 84, 981-991.	0.8	125
51	Molecular Evolution and Diversity of Lignin Degrading Heme Peroxidases in the Agaricomycetes. <i>Journal of Molecular Evolution</i> , 2008, 66, 243-257.	0.8	120
52	After the gold rush, or before the flood? Evolutionary morphology of mushroom-forming fungi (Agaricomycetes) in the early 21st century. <i>Mycological Research</i> , 2007, 111, 1001-1018.	2.5	116
53	Transcriptomic atlas of mushroom development reveals conserved genes behind complex multicellularity in fungi. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 7409-7418.	3.3	115
54	Research Coordination Networks: a phylogeny for kingdom Fungi (Deep Hypha). <i>Mycologia</i> , 2006, 98, 829-837.	0.8	114

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55	Phylogenetic classification of <i>Trametes</i> (Basidiomycota, Polyporales) based on a fiveâ€‘marker dataset. <i>Taxon</i> , 2011, 60, 1567-1583.	0.4	111
56	Genomewide analysis of polysaccharides degrading enzymes in 11 white- and brown-rot Polyporales provides insight into mechanisms of wood decay. <i>Mycologia</i> , 2013, 105, 1412-1427.	0.8	110
57	Fungal ecology catches fire. <i>New Phytologist</i> , 2009, 184, 279-282.	3.5	104
58	Evolutionary dynamics of host specialization in wood-decay fungi. <i>BMC Evolutionary Biology</i> , 2018, 18, 119.	3.2	104
59	The genome of the xerotolerant mold <i>Wallemia sebi</i> reveals adaptations to osmotic stress and suggests cryptic sexual reproduction. <i>Fungal Genetics and Biology</i> , 2012, 49, 217-226.	0.9	103
60	Phylogenetic Relationships of <i>Lentinus</i> (Basidiomycotina) Inferred from Molecular and Morphological Characters. <i>Systematic Botany</i> , 1993, 18, 409.	0.2	101
61	Research Coordination Networks: a phylogeny for kingdom Fungi (Deep Hypha). <i>Mycologia</i> , 2006, 98, 829-837.	0.8	97
62	Trends in Morphological Evolution in Homobasidiomycetes Inferred Using Maximum Likelihood: A Comparison of Binary and Multistate Approaches. <i>Systematic Biology</i> , 2004, 53, 889-903.	2.7	92
63	Phylogenetic relationships of cyphelloid homobasidiomycetes. <i>Molecular Phylogenetics and Evolution</i> , 2004, 33, 501-515.	1.2	92
64	Resolving the phylogenetic position of the Wallemiomycetes: an enigmatic major lineage of Basidiomycota. <i>Canadian Journal of Botany</i> , 2006, 84, 1794-1805.	1.2	91
65	A phylogenetic overview of the Agaricomycotina. <i>Mycologia</i> , 2006, 98, 917-925.	0.8	87
66	Phylogeny of the Pluteaceae (Agaricales, Basidiomycota): taxonomy and character evolution. <i>Fungal Biology</i> , 2011, 115, 1-20.	1.1	86
67	A phylogenetic overview of the antrodia clade (Basidiomycota, Polyporales). <i>Mycologia</i> , 2013, 105, 1391-1411.	0.8	86
68	Revisiting the taxonomy of <i>Phanerochaete</i> (Polyporales, Basidiomycota) using a four gene dataset and extensive ITS sampling. <i>Fungal Biology</i> , 2015, 119, 679-719.	1.1	86
69	Phylogenetic relationships of cantharelloid and clavarioid Homobasidiomycetes based on mitochondrial and nuclear rDNA sequences. <i>Mycologia</i> , 1999, 91, 944-963.	0.8	85
70	An overview of the higher level classification of Pucciniomycotina based on combined analyses of nuclear large and small subunit rDNA sequences. <i>Mycologia</i> , 2006, 98, 896-905.	0.8	80
71	Diversity and evolution of ectomycorrhizal host associations in the Sclerodermatineae (Boletales). <i>Trends in Microbiology</i> , 2014, 22, 314-321.	3.5	73
72	Unexpected diversity of basidiomycetous endophytes in sapwood and leaves of <i>Hevea</i> . <i>Mycologia</i> , 2015, 107, 284-297.	0.8	73

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73	The genome of <i>Xylona heveae</i> provides a window into fungal endophytism. <i>Fungal Biology</i> , 2016, 120, 26-42.	1.1	72
74	Genomic Analysis Enlightens Agaricales Lifestyle Evolution and Increasing Peroxidase Diversity. <i>Molecular Biology and Evolution</i> , 2021, 38, 1428-1446.	3.5	72
75	Molecular phylogenetics of the Gloeophyllales and relative ages of clades of Agaricomycotina producing a brown rot. <i>Mycologia</i> , 2011, 103, 510-524.	0.8	69
76	Genetic Bases of Fungal White Rot Wood Decay Predicted by Phylogenomic Analysis of Correlated Gene-Phenotype Evolution. <i>Molecular Biology and Evolution</i> , 2017, 34, 35-44.	3.5	65
77	Fruiting body form, not nutritional mode, is the major driver of diversification in mushroom-forming fungi. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 32528-32534.	3.3	65
78	Phylogenetic Relationships of Cantharelloid and Clavarioid Homobasidiomycetes Based on Mitochondrial and Nuclear rDNA Sequences. <i>Mycologia</i> , 1999, 91, 944.	0.8	64
79	Toward a phylogenetic classification of the Leotiomycetes based on rDNA data. <i>Mycologia</i> , 2006, 98, 1065-1075.	0.8	64
80	Automated Phylogenetic Taxonomy: An Example in the Homobasidiomycetes (Mushroom-Forming) <i>Trends in Microbiology</i> , 2007, 15, 100-105.	2.7	63
81	EFFECTS OF GASTEROID FRUITING BODY MORPHOLOGY ON DIVERSIFICATION RATES IN THREE INDEPENDENT CLADES OF FUNGI ESTIMATED USING BINARY STATE SPECIATION AND EXTINCTION ANALYSIS. <i>Evolution; International Journal of Organic Evolution</i> , 2011, 65, 1305-1322.	1.1	63
82	Integrating phylogenetic analysis and classification in fungi. <i>Mycologia</i> , 1998, 90, 347-356.	0.8	62
83	Convergent evolution of sequestrate forms in <i>Amanita</i> under Mediterranean climate conditions. <i>Mycologia</i> , 2010, 102, 675-688.	0.8	59
84	The invisible dimension of fungal diversity. <i>Science</i> , 2016, 351, 1150-1151.	6.0	59
85	Evolutionary relationships of <i>Mycaureola dilseae</i> (Agaricales), a basidiomycete pathogen of a subtidal rhodophyte. <i>American Journal of Botany</i> , 2006, 93, 547-556.	0.8	58
86	Polypores and genus concepts in Phanerochaetaceae (Polyporales, Basidiomycota). <i>MycKeys</i> , 0, 17, 1-46.	0.8	54
87	Genomics and Development of <i>Lentinus tigrinus</i> : A White-Rot Wood-Decaying Mushroom with Dimorphic Fruiting Bodies. <i>Genome Biology and Evolution</i> , 2018, 10, 3250-3261.	1.1	53
88	Evolutionary Relationships of <i>Lentinus</i> to the Polyporaceae: Evidence from Restriction Analysis of Enzymatically Amplified Ribosomal Dna. <i>Mycologia</i> , 1991, 83, 425-439.	0.8	52
89	Phylogenetic relationships of <i>Sparassis</i> inferred from nuclear and mitochondrial ribosomal DNA and RNA polymerase sequences. <i>Mycologia</i> , 2004, 96, 1015-1029.	0.8	48
90	Degradation of Bunker C Fuel Oil by White-Rot Fungi in Sawdust Cultures Suggests Potential Applications in Bioremediation. <i>PLoS ONE</i> , 2015, 10, e0130381.	1.1	43

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91	Evolution of Marine Mushrooms. <i>Biological Bulletin</i> , 2001, 201, 319-322.	0.7	42
92	(308â€“310) Proposals to permit DNA sequence data to serve as types of names of fungi. <i>Taxon</i> , 2016, 65, 899-900.	0.4	42
93	Toward genome-enabled mycology. <i>Mycologia</i> , 2013, 105, 1339-1349.	0.8	38
94	Mycophagous rove beetles highlight diverse mushrooms in the Cretaceous. <i>Nature Communications</i> , 2017, 8, 14894.	5.8	38
95	The phylogeny of selected <i>Phylloporus</i> species, inferred from NUC-LSU and ITS sequences, and descriptions of new species from the Old World. <i>Fungal Diversity</i> , 2012, 55, 109-123.	4.7	37
96	Evolutionary Relationships of <i>Lentinus</i> to the Polyporaceae: Evidence from Restriction Analysis of Enzymatically Amplified Ribosomal DNA. <i>Mycologia</i> , 1991, 83, 425.	0.8	36
97	Phylogenetic Relationships of <i>Sparassis</i> Inferred from Nuclear and Mitochondrial Ribosomal DNA and RNA Polymerase Sequences. <i>Mycologia</i> , 2004, 96, 1015.	0.8	35
98	Integrating Phylogenetic Analysis and Classification in Fungi. <i>Mycologia</i> , 1998, 90, 347.	0.8	33
99	Species delimitation in <i>Trametes</i> : a comparison of ITS, RPB1, RPB2 and TEF1 gene phylogenies. <i>Mycologia</i> , 2014, 106, 735-745.	0.8	33
100	Molecular phylogeny and phylogeography of Holarctic species of <i>Pluteus</i> section <i>Pluteus</i> (Agaricales): Tj ETQq0 0 0 rgBT /Overlock 10 Tf	0.5	33
101	Characterization of Three <i>mnp</i> Genes of <i>Fomitiporia mediterranea</i> and Report of Additional Class II Peroxidases in the Order Hymenochaetales. <i>Applied and Environmental Microbiology</i> , 2010, 76, 6431-6440.	1.4	32
102	Phylogenetic relationships and morphological evolution in <i>Lentinus</i> , <i>Polyporellus</i> and <i>Neofavolus</i> , emphasizing southeastern Asian taxa. <i>Mycologia</i> , 2015, 107, 460-474.	0.8	31
103	Phylogenetic relationships of the marine gasteromycete <i>Nia vibrissa</i> . <i>Mycologia</i> , 2001, 93, 679-688.	0.8	29
104	Phylogenetic analyses of <i>Aleurodiscus</i> s.l. and allied genera. <i>Mycologia</i> , 2001, 93, 720-731.	0.8	28
105	Another fossil agaric from Dominican amber. <i>Mycologia</i> , 2003, 95, 685-687.	0.8	27
106	Phylogenetic taxon definitions for Fungi, Dikarya, Ascomycota and Basidiomycota. <i>IMA Fungus</i> , 2018, 9, 291-298.	1.7	26
107	Phylogeny and a new species of <i>Sparassis</i> (Polyporales, Basidiomycota): evidence from mitochondrial <i>atp6</i> , nuclear <i>rDNA</i> and <i>rpb2</i> genes. <i>Mycologia</i> , 2006, 98, 584-592.	0.8	25
108	Climate, decay, and the death of the coal forests. <i>Current Biology</i> , 2016, 26, R563-R567.	1.8	25

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109	Where are all the undocumented fungal species? A study of <i>Mortierella</i> demonstrates the need for sequence-based classification. <i>New Phytologist</i> , 2011, 191, 592-596.	3.5	24
110	Substrate-Specific Differential Gene Expression and RNA Editing in the Brown Rot Fungus <i>Fomitopsis pinicola</i> . <i>Applied and Environmental Microbiology</i> , 2018, 84, .	1.4	22
111	Towards the unification of sequence-based classification and sequence-based identification of host-associated microorganisms. <i>New Phytologist</i> , 2015, 205, 27-31.	3.5	21
112	Evolutionary transition to the ectomycorrhizal habit in the genomes of a hyperdiverse lineage of mushroom-forming fungi. <i>New Phytologist</i> , 2022, 233, 2294-2309.	3.5	21
113	Phylogenetic Relationships of the Marine Gasteromycete <i>Nia vibrissa</i> . <i>Mycologia</i> , 2001, 93, 679.	0.8	19
114	Draft genome sequence of a monokaryotic model brown-rot fungus <i>Postia</i> ( <i>Rhodonia</i> ) <i>placenta</i> SB12. <i>Genomics Data</i> , 2017, 14, 21-23.	1.3	19
115	The Secotioid Form of <i>Lentinus tigrinus</i> : Genetics and Development of a Fungal Morphological Innovation. <i>American Journal of Botany</i> , 1994, 81, 466.	0.8	18
116	Hymenophore Development and Evolution in <i>Lentinus</i> . <i>Mycologia</i> , 1993, 85, 428-443.	0.8	17
117	<i>Sparassis cystidiosa</i> sp. nov. from Thailand is described using morphological and molecular data. <i>Mycologia</i> , 2004, 96, 1010-1014.	0.8	16
118	Population genomics provides insights into the genetic basis of adaptive evolution in the mushroom-forming fungus <i>Lentinula edodes</i> . <i>Journal of Advanced Research</i> , 2022, 38, 91-106.	4.4	16
119	Draft Genome Sequence of the White-Rot Fungus <i>Obba rivulosa</i> 3A-2. <i>Genome Announcements</i> , 2016, 4, .	0.8	15
120	<i>Neocampanella</i> , a new corticioid fungal genus, and a note on <i>Dendrothele bispora</i> . <i>Botany</i> , 2009, 87, 875-882.	0.5	13
121	Evolutionary Morphogenesis of Sexual Fruiting Bodies in Basidiomycota: Toward a New Evo-Devo Synthesis. <i>Microbiology and Molecular Biology Reviews</i> , 2022, 86, e0001921.	2.9	13
122	Hymenophore Development and Evolution in <i>Lentinus</i> . <i>Mycologia</i> , 1993, 85, 428.	0.8	12
123	Phylogeny and genetic diversity of <i>Bridgeoporus nobilissimus</i> inferred using mitochondrial and nuclear rDNA sequences. <i>Mycologia</i> , 2003, 95, 836-845.	0.8	12
124	Proposals for consideration at IMC11 to modify provisions related solely to fungi in the International Code of Nomenclature for algae, fungi, and plants. <i>IMA Fungus</i> , 2018, 9, i-vii.	1.7	10
125	Sporocarp ontogeny in <i>Panus</i> (Basidiomycotina): evolution and classification. <i>American Journal of Botany</i> , 1993, 80, 1336-1348.	0.8	9
126	Global phylogeny of the Shiitake mushroom and related <i>Lentinula</i> species uncovers novel diversity and suggests an origin in the Neotropics. <i>Molecular Phylogenetics and Evolution</i> , 2022, 173, 107494.	1.2	8



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127	Toward Sequence-Based Classification of Fungal Species. <i>IMA Fungus</i> , 2013, 4, A33-A34.	1.7	7
128	Sporocarp ontogeny in <i>Panus</i> (Basidiomycotina): evolution and classification. , 1993, 80, 1336.		6
129	RNA-editing in Basidiomycota, revisited. <i>ISME Communications</i> , 2021, 1, .	1.7	2
130	Digital identifiers for fungal speciesâ€™Response. <i>Science</i> , 2016, 352, 1183-1183.	6.0	1
131	ll.14. Major Events in the Evolution of Fungi. , 2013, , 152-158.		0