

Irene Barnes

List of Publications by Year in descending order

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Version: 2024-02-01

260
papers

12,101
citations

29994

54
h-index

39575

94
g-index

261
all docs

261
docs citations

261
times ranked

8030
citing authors

#	ARTICLE	IF	CITATIONS
1	<i>Lecanosticta pharomachri</i> and Its Newly Discovered Sexual State Causing a Serious Needle Disease of <i>Pinus</i> spp. in Colombia. <i>Plant Disease</i> , 2022, 106, 1935-1943.	0.7	4
2	<i>Calonectria</i> species, including four novel taxa, associated with Eucalyptus in Malaysia. <i>Mycological Progress</i> , 2022, 21, 181-197.	0.5	11
3	IMA Genome - F16. <i>IMA Fungus</i> , 2022, 13, 3.	1.7	4
4	Invasion Frameworks: a Forest Pathogen Perspective. <i>Current Forestry Reports</i> , 2022, 8, 74-89.	3.4	14
5	<i>Calonectria</i> in the age of genes and genomes: Towards understanding an important but relatively unknown group of pathogens. <i>Molecular Plant Pathology</i> , 2022, 23, 1060-1072.	2.0	9
6	<i>Botryosphaeriaceae</i> diversity on <i>Eucalyptus</i> clones in different climate zones of Indonesia. <i>Forest Pathology</i> , 2022, 52, .	0.5	4
7	Novel mating-type-associated genes and gene fragments in the genomes of <i>Mycosphaerellaceae</i> and <i>Teratosphaeriaceae</i> fungi. <i>Molecular Phylogenetics and Evolution</i> , 2022, 171, 107456.	1.2	0
8	Pathogens of the <i>Araucariaceae</i> : How Much Do We Know?. <i>Current Forestry Reports</i> , 2022, 8, 124-147.	3.4	3
9	<i>Dothistroma</i> needle blight. , 2022, , 179-199.		7
10	Phylogenomic Analysis of a 55.1-kb 19-Gene Dataset Resolves a Monophyletic <i>Fusarium</i> that Includes the <i>Fusarium solani</i> Species Complex. <i>Phytopathology</i> , 2021, 111, 1064-1079.	1.1	107
11	Susceptibility of native New Zealand Myrtaceae to the South African strain of <i>Austropuccinia psidii</i> : A biosecurity threat. <i>Plant Pathology</i> , 2021, 70, 667-675.	1.2	6
12	Worldwide Genetic Structure Elucidates the Eurasian Origin and Invasion Pathways of <i>Dothistroma septosporum</i> , Causal Agent of <i>Dothistroma</i> Needle Blight. <i>Journal of Fungi (Basel, Switzerland)</i> , 2021, 7, 111.	1.5	14
13	<i>Eucalyptus</i> scab and shoot malformation: A new and serious foliar disease of <i>Eucalyptus</i> caused by <i>Elsinoe necatrix</i> sp. nov.. <i>Plant Pathology</i> , 2021, 70, 1230-1242.	1.2	11
14	<i>Pewenomyces kutranfy</i> gen. nov. et sp. nov. causal agent of an important canker disease on <i>Araucaria araucana</i> in Chile. <i>Plant Pathology</i> , 2021, 70, 1243-1259.	1.2	8
15	Ophiostomatalean fungi associated with wood boring beetles in South Africa including two new species. <i>Antonie Van Leeuwenhoek</i> , 2021, 114, 667-686.	0.7	7
16	Doing it alone: Unisexual reproduction in filamentous ascomycete fungi. <i>Fungal Biology Reviews</i> , 2021, 35, 1-13.	1.9	20
17	Genetic recombination in <i>Teratosphaeria destructans</i> causing a new disease outbreak in Malaysia. <i>Forest Pathology</i> , 2021, 51, e12683.	0.5	9
18	Novel <i>Fusarium</i> mutualists of two <i>Euwallacea</i> species infesting <i>Acacia crassicarpa</i> in Indonesia. <i>Mycologia</i> , 2021, 113, 536-558.	0.8	9

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19	Rapid Detection of Pine Pathogens <i>Lecanosticta acicola</i> , <i>Dothistroma pini</i> and <i>D. septosporum</i> on Needles by Probe-Based LAMP Assays. <i>Forests</i> , 2021, 12, 479.	0.9	15
20	Population genetic structure and migration patterns of the maize pathogenic fungus, <i>Cercospora zeina</i> in East and Southern Africa. <i>Fungal Genetics and Biology</i> , 2021, 149, 103527.	0.9	7
21	Botanical gardens as key resources and hazards for biosecurity. <i>Biodiversity and Conservation</i> , 2021, 30, 1929-1946.	1.2	21
22	Global Genetic Diversity and Mating Type Distribution of <i>Calonectria pauciramosa</i> : An Important Wide-Host-Range Plant Pathogen. <i>Plant Disease</i> , 2021, 105, 1648-1656.	0.7	6
23	Population structure and genetic diversity suggest recent introductions of <i>Dothistroma pini</i> in Slovakia. <i>Plant Pathology</i> , 2021, 70, 1883-1896.	1.2	5
24	Spatial Genetic Structure of the Insect-Vectored Conifer Pathogen <i>Leptographium wageneri</i> Suggests Long Distance Gene Flow Among Douglas-fir Plantations in Western Oregon. <i>Frontiers in Forests and Global Change</i> , 2021, 4, .	1.0	1
25	Genetic Networks That Govern Sexual Reproduction in the Pezizomycotina. <i>Microbiology and Molecular Biology Reviews</i> , 2021, 85, e0002021.	2.9	14
26	<i>Ceratocystis ficicola</i> causing a serious disease of <i>Ficus carica</i> in Greece. <i>Phytopathologia Mediterranea</i> , 2021, 60, 337-349.	0.6	7
27	Population Diversity and Genetic Structure Reveal Patterns of Host Association and Anthropogenic Impact for the Globally Important Fungal Tree Pathogen <i>Ceratocystis manginecans</i> . <i>Journal of Fungi (Basel, Switzerland)</i> , 2021, 7, 759.	1.5	4
28	Signatures of Post-Glacial Genetic Isolation and Human-Driven Migration in the <i>Dothistroma</i> Needle Blight Pathogen in Western Canada. <i>Phytopathology</i> , 2021, 111, 116-127.	1.1	15
29	New hosts for <i>Lecanosticta acicola</i> and <i>Dothistroma septosporum</i> in newly established arboreta in Spain. <i>Forest Pathology</i> , 2021, 51, .	0.5	5
30	Sexual reproduction in populations of <i>Austropuccinia psidii</i> . <i>European Journal of Plant Pathology</i> , 2020, 156, 537-545.	0.8	8
31	Highly transferable microsatellite markers for the genera <i>Lasiodiplodia</i> and <i>Neofusicoccum</i> . <i>Fungal Ecology</i> , 2020, 44, 100903.	0.7	3
32	<i>DsEcp2-1</i> is a polymorphic effector that restricts growth of <i>Dothistroma septosporum</i> in pine. <i>Fungal Genetics and Biology</i> , 2020, 135, 103300.	0.9	14
33	Pathogenicity of <i>Chrysoporthe deuterocubensis</i> and <i>MyrtoPORthe bodenii</i> gen. et sp. nov. on <i>Eucalyptus</i> in Sabah, Malaysia. <i>Australasian Plant Pathology</i> , 2020, 49, 53-64.	0.5	11
34	<i>Ceratocystis quercicola</i> sp. nov. from <i>Quercus variabilis</i> in Korea. <i>Mycobiology</i> , 2020, 48, 245-251.	0.6	2
35	Mating strategy and mating type distribution in six global populations of the <i>Eucalyptus</i> foliar pathogen <i>Teratosphaeria destructans</i> . <i>Fungal Genetics and Biology</i> , 2020, 137, 103350.	0.9	19
36	Epitypification of <i>Ceratocystis fimbriata</i> . <i>Fungal Systematics and Evolution</i> , 2020, 6, 289-298.	0.9	9

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37	Phylogenomic incongruence in <i>Ceratocystis</i> : a clue to speciation?. <i>BMC Genomics</i> , 2020, 21, 362.	1.2	11
38	<i>Ceratocystis</i> wilt on <i>Eucalyptus</i> : first record from South Africa. <i>Southern Forests</i> , 2020, 82, 24-31.	0.2	9
39	Bark beetle mycobiome: collaboratively defined research priorities on a widespread insect-fungus symbiosis. <i>Symbiosis</i> , 2020, 81, 101-113.	1.2	20
40	Low genetic diversity and strong geographic structure in introduced populations of the <i>Eucalyptus</i> foliar pathogen <i>Teratosphaeria destructans</i> . <i>Plant Pathology</i> , 2020, 69, 1540-1550.	1.2	9
41	Genome comparisons suggest an association between <i>Ceratocystis</i> host adaptations and effector clusters in unique transposable element families. <i>Fungal Genetics and Biology</i> , 2020, 143, 103433.	0.9	9
42	<i>Euwallacea perbrevis</i> (Coleoptera: Curculionidae: Scolytinae), a confirmed pest on <i>Acacia crassicarpa</i> in Riau, Indonesia, and a new fungal symbiont; <i>Fusarium rekanum</i> sp. nov.. <i>Antonie Van Leeuwenhoek</i> , 2020, 113, 803-823.	0.7	21
43	Quantification of Outcrossing Events in Haploid Fungi Using Microsatellite Markers. <i>Journal of Fungi</i> (Basel, Switzerland), 2020, 6, 48.	1.5	1
44	Tree health in South Africa: Retrospect and prospect. <i>South African Journal of Science</i> , 2020, 116, .	0.3	3
45	Mating genes in <i>Calonectria</i> and evidence for a heterothallic ancestral state. <i>Persoonia: Molecular Phylogeny and Evolution of Fungi</i> , 2020, 45, 163-176.	1.6	20
46	<i>Caliciopsis moriondi</i> , a new species for a fungus long confused with the pine pathogen <i>C. pinea</i> . <i>MycKeys</i> , 2020, 73, 87-108.	0.8	7
47	<i>Cryphonectriaceae</i> associated with rust-infected <i>Syzygium jambos</i> in Hawaii. <i>MycKeys</i> , 2020, 76, 49-79.	0.8	9
48	Novel species of <i>Huntia</i> from naturally-occurring forest trees in Greece and South Africa. <i>MycKeys</i> , 2020, 69, 33-52.	0.8	3
49	<i>Lecanosticta acicola</i> : A growing threat to expanding global pine forests and plantations. <i>Molecular Plant Pathology</i> , 2019, 20, 1327-1364.	2.0	28
50	QTL mapping of mycelial growth and aggressiveness to distinct hosts in <i>Ceratocystis</i> pathogens. <i>Fungal Genetics and Biology</i> , 2019, 131, 103242.	0.9	12
51	Influence of farming practices on the population genetics of the maize pathogen <i>Cercospora zeina</i> in South Africa. <i>Fungal Genetics and Biology</i> , 2019, 125, 36-44.	0.9	13
52	Biodiversity of <i>Lecanosticta</i> pine-needle blight pathogens suggests a Mesoamerican Centre of origin. <i>IMA Fungus</i> , 2019, 10, 2.	1.7	19
53	Global population genomics of the forest pathogen <i>Dothistroma septosporum</i> reveal chromosome duplications in high dothistromin-producing strains. <i>Molecular Plant Pathology</i> , 2019, 20, 784-799.	2.0	19
54	It's All in the Genes: The Regulatory Pathways of Sexual Reproduction in Filamentous Ascomycetes. <i>Genes</i> , 2019, 10, 330.	1.0	31

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55	Genetic diversity of the pine pathogen <i>Lecanosticta acicola</i> in Slovenia and Croatia. <i>Plant Pathology</i> , 2019, 68, 1120-1131.	1.2	12
56	The pandemic strain of <i>Austropuccinia psidii</i> causes myrtle rust in New Zealand and Singapore. <i>Australasian Plant Pathology</i> , 2019, 48, 253-256.	0.5	14
57	Genomic analysis of the aggressive tree pathogen <i>Ceratocystis albifundus</i> . <i>Fungal Biology</i> , 2019, 123, 351-363.	1.1	11
58	IMA Genome-F 11. <i>IMA Fungus</i> , 2019, 10, 13.	1.7	12
59	Draft genome sequences of five <i>Calonectria</i> species from Eucalyptus plantations in China, <i>Celoporthe dispersa</i> , <i>Sporothrix phasma</i> and <i>Alectoria sarmentosa</i> . <i>IMA Fungus</i> , 2019, 10, 22.	1.7	17
60	Ten new species of <i>Calonectria</i> from Indonesia and Vietnam. <i>Mycologia</i> , 2019, 111, 78-102.	0.8	38
61	Teratosphaeria stem canker of <i>Eucalyptus</i> : two pathogens, one devastating disease. <i>Molecular Plant Pathology</i> , 2019, 20, 8-19.	2.0	37
62	Inheritance of phenotypic traits in the progeny of a <i>Ceratocystis</i> interspecific cross. <i>Fungal Biology</i> , 2018, 122, 717-729.	1.1	6
63	Looking for relationships between the populations of <i>Dothistroma septosporum</i> in northern Europe and Asia. <i>Fungal Genetics and Biology</i> , 2018, 110, 15-25.	0.9	23
64	Non-Mendelian segregation influences the infection biology and genetic structure of the African tree pathogen <i>Ceratocystis albifundus</i> . <i>Fungal Biology</i> , 2018, 122, 222-230.	1.1	4
65	Molecular phylogenetics and microsatellite analysis reveal a new pathogenic <i>Ceratocystis</i> species in the Asian–Australian clade. <i>Plant Pathology</i> , 2018, 67, 1097-1113.	1.2	14
66	Evidence that <i>Austropuccinia psidii</i> may complete its sexual life cycle on Myrtaceae. <i>Plant Pathology</i> , 2018, 67, 729-734.	1.2	19
67	New species of <i>Cylindrocladiella</i> from plantation soils in South-East Asia. <i>MycKeys</i> , 2018, 32, 1-24.	0.8	1
68	Nine draft genome sequences of <i>Claviceps purpurea</i> s.lat., including <i>C. arundinis</i> , <i>C. humidiphila</i> , and <i>C. cf. spartinae</i> , pseudomolecules for the pitch canker pathogen <i>Fusarium circinatum</i> , draft genome of <i>Davidsoniella eucalypti</i> , <i>Grosmannia galeiformis</i> , <i>Quambalaria eucalypti</i> , and <i>Teratosphaeria destructans</i> . <i>IMA Fungus</i> , 2018, 9, 401-418.	1.7	31
69	Nine novel species of <i>Huntiella</i> from southern China with three distinct mating strategies and variable levels of pathogenicity. <i>Mycologia</i> , 2018, 110, 1145-1171.	0.8	7
70	Armillaria Root-Rot Pathogens: Species Boundaries and Global Distribution. <i>Pathogens</i> , 2018, 7, 83.	1.2	40
71	<i>Ceratocystidaceae</i> exhibit high levels of recombination at the mating-type (MAT) locus. <i>Fungal Biology</i> , 2018, 122, 1184-1191.	1.1	10
72	New <i>Ceratocystis</i> species associated with rapid death of <i>Metrosideros polymorpha</i> in Hawai'i. <i>Persoonia: Molecular Phylogeny and Evolution of Fungi</i> , 2018, 40, 154-181.	1.6	106

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73	Draft genome sequence of <i>Annulohyphomyces stylum</i> , <i>Aspergillus mulundensis</i> , <i>Berkeleyomyces basicola</i> (syn. <i>Thielaviopsis basicola</i>), <i>Ceratocystis smalleyi</i> , two <i>Cercospora beticola</i> strains, <i>Coleophoma cylindrospora</i> , <i>Fusarium fracticaudum</i> , <i>Phialophora</i> cf. <i>hyalina</i> , and <i>Morchella septimelata</i> . <i>IMA Fungus</i> , 2018, 9, 199-223.	1.7	37
74	Genetic diversity of <i>Amylostereum areolatum</i> , the fungal symbiont of the invasive woodwasp <i>Sirex noctilio</i> in South Africa. <i>Forest Pathology</i> , 2018, 48, e12449.	0.5	6
75	Canker Stain: A Lethal Disease Destroying Iconic Plane Trees. <i>Plant Disease</i> , 2017, 101, 645-658.	0.7	66
76	Genera of phytopathogenic fungi: GOPHY 1. <i>Studies in Mycology</i> , 2017, 86, 99-216.	4.5	276
77	The pandemic biotype of <i>Austropuccinia psidii</i> discovered in South America. <i>Australasian Plant Pathology</i> , 2017, 46, 267-275.	0.5	18
78	The unified framework for biological invasions: a forest fungal pathogen perspective. <i>Biological Invasions</i> , 2017, 19, 3201-3214.	1.2	35
79	Novel associations between ophiostomatoid fungi, insects and tree hosts: current status and future prospects. <i>Biological Invasions</i> , 2017, 19, 3215-3228.	1.2	49
80	Pathogens on the Move: A 100-Year Global Experiment with Planted Eucalypts. <i>BioScience</i> , 2017, 67, 14-25.	2.2	96
81	Urban trees: bridge-heads for forest pest invasions and sentinels for early detection. <i>Biological Invasions</i> , 2017, 19, 3515-3526.	1.2	89
82	Biological invasions in forest ecosystems. <i>Biological Invasions</i> , 2017, 19, 3437-3458.	1.2	161
83	Evolution of polyketide synthesis in a Dothideomycete forest pathogen. <i>Fungal Genetics and Biology</i> , 2017, 106, 42-50.	0.9	8
84	<i>Botryosphaeria dothidea</i> : a latent pathogen of global importance to woody plant health. <i>Molecular Plant Pathology</i> , 2017, 18, 477-488.	2.0	202
85	Ecology and population structure of a tree wound-infecting fungus in a native South African forest environment. <i>Fungal Biology</i> , 2017, 121, 69-81.	1.1	5
86	Putative origins of the fungus <i>Leptographium procerum</i> . <i>Fungal Biology</i> , 2017, 121, 82-94.	1.1	12
87	Invasive Everywhere? Phylogeographic Analysis of the Globally Distributed Tree Pathogen <i>Lasiodiplodia theobromae</i> . <i>Forests</i> , 2017, 8, 145.	0.9	31
88	Draft genome of <i>Cercospora zeina</i> , <i>Fusarium pininemorale</i> , <i>Hawksworthiomyces lignivorus</i> , <i>Huntia decipiens</i> and <i>Ophiostoma ips</i> . <i>IMA Fungus</i> , 2017, 8, 385-396.	1.7	37
89	A plant pathology perspective of fungal genome sequencing. <i>IMA Fungus</i> , 2017, 8, 1-15.	1.7	75
90	<i>Calonectria</i> species isolated from Eucalyptus plantations and nurseries in South China. <i>IMA Fungus</i> , 2017, 8, 259-286.	1.7	37

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91	Ophiostomatoid fungi associated with conifer-infesting beetles and their phoretic mites in Yunnan, China. <i>MycKeys</i> , 2017, 28, 19-64.	0.8	43
92	Draft genome sequences for <i>Ceratocystis fagacearum</i> , <i>C. harringtonii</i> , <i>Grosmannia penicillata</i> , and <i>Huntia bhutanensis</i> . <i>IMA Fungus</i> , 2016, 7, 317-323.	1.7	31
93	Fungal Genomics Challenges the Dogma of Name-Based Biosecurity. <i>PLoS Pathogens</i> , 2016, 12, e1005475.	2.1	36
94	Ecological disequilibrium drives insect pest and pathogen accumulation in non-native trees. <i>AoB PLANTS</i> , 2016, , plw081.	1.2	25
95	Lack of Population Structure and Mixed Reproduction Modes in <i>Exserohilum turcicum</i> from South Africa. <i>Phytopathology</i> , 2016, 106, 1386-1392.	1.1	15
96	<i>Cercospora zeina</i> from Maize in South Africa Exhibits High Genetic Diversity and Lack of Regional Population Differentiation. <i>Phytopathology</i> , 2016, 106, 1194-1205.	1.1	11
97	Endophytic Cryphonectriaceae on native Myrtales: Possible origin of <i>Chrysoporthe</i> canker on plantation-grown Eucalyptus. <i>Fungal Biology</i> , 2016, 120, 827-835.	1.1	12
98	The Eucalyptus shoot and leaf pathogen <i>Teratosphaeria destructans</i> recorded in South Africa. <i>Southern Forests</i> , 2016, 78, 123-129.	0.2	18
99	Population structure of <i>Holocryphia capensis</i> (cryphonectriaceae) from <i>Metrosideros angustifolia</i> and its pathogenicity to Eucalyptus species. <i>Australasian Plant Pathology</i> , 2016, 45, 201-207.	0.5	4
100	Three genetic groups of the Eucalyptus stem canker pathogen <i>Teratosphaeria zuluensis</i> introduced into Africa from an unknown source. <i>Antonie Van Leeuwenhoek</i> , 2016, 109, 21-33.	0.7	6
101	Neotypification of <i>Dothistroma septosporum</i> and epitypification of <i>D. Æpini</i> causal agents of <i>Dothistroma</i> needle blight of pine. <i>Forest Pathology</i> , 2016, 46, 388-407.	0.5	38
102	A worldwide perspective on the management and control of <i>Dothistroma</i> needle blight. <i>Forest Pathology</i> , 2016, 46, 472-488.	0.5	58
103	<i>Dothistroma</i> needle blight: an emerging epidemic caused by <i>Dothistroma septosporum</i> in Colombia. <i>Plant Pathology</i> , 2016, 65, 53-63.	1.2	25
104	Genetic Analyses Suggest Separate Introductions of the Pine Pathogen <i>Lecanosticta acicola</i> Into Europe. <i>Phytopathology</i> , 2016, 106, 1413-1425.	1.1	30
105	Global food and fibre security threatened by current inefficiencies in fungal identification. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2016, 371, 20160024.	1.8	74
106	Development of microsatellite markers for the pine needle blight pathogen, <i>Dothistroma pini</i> . <i>Forest Pathology</i> , 2016, 46, 497-506.	0.5	11
107	Global geographic distribution and host range of <i>Dothistroma</i> species: a comprehensive review. <i>Forest Pathology</i> , 2016, 46, 408-442.	0.5	84
108	A unique genotype of the rust pathogen, <i>Puccinia psidii</i> , on Myrtaceae in South Africa. <i>Australasian Plant Pathology</i> , 2016, 45, 645-652.	0.5	32

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109	A review of Pinaceae resistance mechanisms against needle and shoot pathogens with a focus on the <i>Dothistroma</i> – <i>Pinus</i> interaction. <i>Forest Pathology</i> , 2016, 46, 453-471.	0.5	26
110	New host range and distribution of <i>Ceratocystis pirilliformis</i> in South Africa. <i>European Journal of Plant Pathology</i> , 2016, 146, 483-496.	0.8	5
111	The genetic landscape of <i>Ceratocystis albifundus</i> populations in South Africa reveals a recent fungal introduction event. <i>Fungal Biology</i> , 2016, 120, 690-700.	1.1	37
112	A possible centre of diversity in South East Asia for the tree pathogen, <i>Ceratocystis manginecans</i> . <i>Infection, Genetics and Evolution</i> , 2016, 41, 73-83.	1.0	25
113	Population genetics and symbiont assemblages support opposing invasion scenarios for the red turpentine beetle (<i>Dendroctonus valens</i>). <i>Biological Journal of the Linnean Society</i> , 2016, 118, 486-502.	0.7	18
114	Novel and co-evolved associations between insects and microorganisms as drivers of forest pestilence. <i>Biological Invasions</i> , 2016, 18, 1045-1056.	1.2	96
115	Exotic biological control agents: A solution or contribution to arthropod invasions?. <i>Biological Invasions</i> , 2016, 18, 953-969.	1.2	131
116	Increasing numbers and intercontinental spread of invasive insects on eucalypts. <i>Biological Invasions</i> , 2016, 18, 921-933.	1.2	134
117	Insects and Diseases of Mediterranean Forests: A South African Perspective. , 2016, , 397-430.		8
118	Population structure and reproductive mode of <i>Dothistroma septosporum</i> in the Brittany peninsula of France. <i>European Journal of Plant Pathology</i> , 2015, 143, 261-275.	0.8	23
119	Homothallism: an umbrella term for describing diverse sexual behaviours. <i>IMA Fungus</i> , 2015, 6, 207-214.	1.7	75
120	Identifying and Naming Plant-Pathogenic Fungi: Past, Present, and Future. <i>Annual Review of Phytopathology</i> , 2015, 53, 247-267.	3.5	115
121	Fungal Planet description sheets: 320–370. <i>Persoonia: Molecular Phylogeny and Evolution of Fungi</i> , 2015, 34, 167-266.	1.6	193
122	Draft genome sequences of <i>Ceratocystis eucalypticola</i> , <i>Chrysosporthe cubensis</i> , <i>C. deuterocubensis</i> , <i>Davidsoniella virescens</i> , <i>Fusarium temperatum</i> , <i>Graphilbum fragrans</i> , <i>Penicillium nordicum</i> , and <i>Thielaviopsis musarum</i> . <i>IMA Fungus</i> , 2015, 6, 493-506.	1.7	57
123	Independent origins and incipient speciation among host-associated populations of <i>Thielaviopsis ethacetica</i> in Cameroon. <i>Fungal Biology</i> , 2015, 119, 957-972.	1.1	5
124	One fungus, which genes? Development and assessment of universal primers for potential secondary fungal DNA barcodes. <i>Persoonia: Molecular Phylogeny and Evolution of Fungi</i> , 2015, 35, 242-263.	1.6	416
125	Molecular markers delimit cryptic species in <i>Ceratocystis sensu stricto</i> . <i>Mycological Progress</i> , 2015, 14, 1.	0.5	47
126	Evaluating the inheritance of <i>Ceratocystis acaciivora</i> symptom expression in a diverse <i>Acacia mangium</i> breeding population. <i>Southern Forests</i> , 2015, 77, 83-90.	0.2	35

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127	New <i>Ceratocystis</i> species from Eucalyptus and Cunninghamia in South China. <i>Antonie Van Leeuwenhoek</i> , 2015, 107, 1451-1473.	0.7	20
128	Changes in planted forests and future global implications. <i>Forest Ecology and Management</i> , 2015, 352, 57-67.	1.4	515
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154	Ion Torrent PGM as Tool for Fungal Community Analysis: A Case Study of Endophytes in <i>Eucalyptus grandis</i> Reveals High Taxonomic Diversity. <i>PLoS ONE</i> , 2013, 8, e81718.	1.1	84
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157	Large Shift in Symbiont Assemblage in the Invasive Red Turpentine Beetle. <i>PLoS ONE</i> , 2013, 8, e78126.	1.1	51
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160	First fungal genome sequence from Africa: A preliminary analysis. <i>South African Journal of Science</i> , 2012, 108, .	0.3	38
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175	Anthropogenic effects on interaction outcomes: examples from insect-microbial symbioses in forest and savanna ecosystems. <i>Symbiosis</i> , 2011, 53, 101-121.	1.2	26
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213	Isolation and characterization of microsatellite loci in <i>Cylindrocladium parasiticum</i> . Molecular Ecology Notes, 2006, 6, 110-112.	1.7	8
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232	<i>Ceratocystis fimbriata</i> infecting <i>Eucalyptus grandis</i> in Uruguay. <i>Australasian Plant Pathology</i> , 2003, 32, 361.	0.5	36
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234	<i>Ceratocystis pirilliformis</i> , a new species from <i>Eucalyptus nitens</i> in Australia. <i>Mycologia</i> , 2003, 95, 865-871.	0.8	46

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254	<i>Kirramyces destructans</i> sp. nov., a serious leaf pathogen of <i>Eucalyptus</i> in Indonesia. <i>South African Journal of Botany</i> , 1996, 62, 325-327.	1.2	36
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257	Comparison of <i>Seiridium</i> Isolates Associated with Cypress Canker Using Sequence Data. <i>Experimental Mycology</i> , 1993, 17, 323-328.	1.8	28
258	<i>Bretziella</i> , a new genus to accommodate the oak wilt fungus, <i>Ceratocystis fagacearum</i> (Microascales), <i>Tj ETQq0 0 0 rgBT /Overlock 10 T</i>	0.8	36
259	Pathologists and entomologists must join forces against forest pest and pathogen invasions. <i>NeoBiota</i> , 0, 58, 107-127.	1.0	28
260	Harmonising the fields of invasion science and forest pathology. <i>NeoBiota</i> , 0, 62, 301-332.	1.0	16