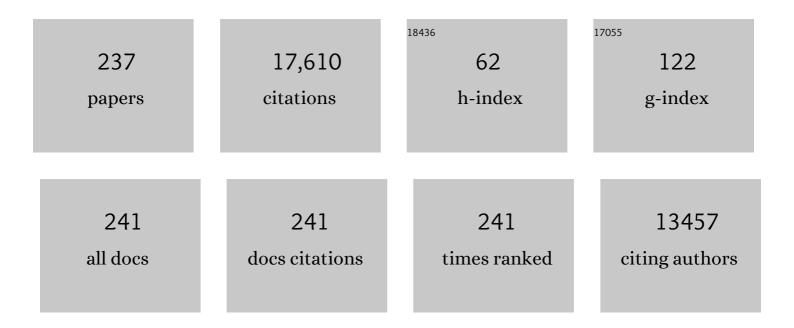
List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Meanderella rijsii, a new opportunist in the fungal order Pleosporales. Microbes and Infection, 2022, 24, 104932.	1.0	1
2	Phylogenetic Revision and Patterns of Host Specificity in the Fungal Subphylum Entomophthoromycotina. Microorganisms, 2022, 10, 256.	1.6	5
3	Black fungi and ants: a genomic comparison of species inhabiting carton nests versus domatia. IMA Fungus, 2022, 13, 4.	1.7	6
4	Phylogenetic and ecological reevaluation of the order Onygenales. Fungal Diversity, 2022, 115, 1-72.	4.7	16
5	Emerging Animal-Associated Fungal Diseases. Journal of Fungi (Basel, Switzerland), 2022, 8, 611.	1.5	11
6	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. Phytopathology, 2021, 111, 1064-1079.	1.1	107
7	Novel black yeast-like species in chaetothyriales with ant-associated life styles. Fungal Biology, 2021, 125, 276-284.	1.1	9
8	Regional Differences in Antifungal Susceptibility of the Prevalent Dermatophyte Trichophyton rubrum. Mycopathologia, 2021, 186, 53-70.	1.3	11
9	Chromoblastomycosis Caused by Phialophora—Proven Cases from Mexico. Journal of Fungi (Basel,) Tj ETQq1	1 0.78431 1.5	4 rgBT /Overle
10	Taxonomy of the Trichophyton mentagrophytes/T. interdigitale Species Complex Harboring the Highly Virulent, Multiresistant Genotype T. indotineae. Mycopathologia, 2021, 186, 315-326.	1.3	76
11	Disseminated Cryptococcosis Presenting as Cellulitis Diagnosed by Laser Capture Microdissection: A Case Report and Literature Review. Mycopathologia, 2021, 186, 423-433.	1.3	4
12	Recent developments in less known and multi-resistant fungal opportunists. Critical Reviews in Microbiology, 2021, 47, 762-780.	2.7	1
13	Majocchi's granuloma: Autoinoculation and adaption of Trichophyton rubrum with molecular evidence. Mycoses, 2021, 64, 1272-1278.	1.8	0
14	Sequence data from isolated lichen-associated melanized fungi enhance delimitation of two new lineages within Chaetothyriomycetidae. Mycological Progress, 2021, 20, 911-927.	0.5	11
15	Molecular and MALDIâ€ToF MS differentiation and antifungal susceptibility of prevalent clinical Fusarium species in China. Mycoses, 2021, 64, 1261-1271.	1.8	7
16	In vitro activities of 8 antifungal drugs against 126 clinical and environmental <i>Exophiala</i> isolates. Mycoses, 2021, 64, 1328-1333.	1.8	3
17	Global guideline for the diagnosis and management of rare mould infections: an initiative of the European Confederation of Medical Mycology in cooperation with the International Society for Human and Animal Mycology and the American Society for Microbiology. Lancet Infectious Diseases, The, 2021, 21, e246-e257.	4.6	167
18	Basidiobolus omanensis sp. nov. Causing Angioinvasive Abdominal Basidiobolomycosis. Journal of Fungi (Basel, Switzerland), 2021, 7, 653.	1.5	7

#	Article	IF	CITATIONS
19	The global burden of chromoblastomycosis. PLoS Neglected Tropical Diseases, 2021, 15, e0009611.	1.3	40
20	Comparative genomics of opportunistic <i>Phialophora</i> species involved in divergent disease types. Mycoses, 2021, 64, 555-568.	1.8	7
21	Development and Analysis of qPCR for the Identification of Arthroconidial Yeasts of the Genus Magnusiomyces. Mycopathologia, 2021, 186, 41-51.	1.3	1
22	Volatile Organic Compounds in the Azteca/Cecropia Ant-Plant Symbiosis and the Role of Black Fungi. Journal of Fungi (Basel, Switzerland), 2021, 7, 836.	1.5	5
23	Molecular Mechanisms of 5-Fluorocytosine Resistance in Yeasts and Filamentous Fungi. Journal of Fungi (Basel, Switzerland), 2021, 7, 909.	1.5	29
24	<i>In vitro</i> activity of eight antifungal drugs against <i>Chaetomiaceae</i> . Medical Mycology, 2021, 60, .	0.3	1
25	Polyphasic Discrimination of Trichophyton tonsurans and T. equinum from Humans and Horses. Mycopathologia, 2020, 185, 113-122.	1.3	19
26	Whole-genome resequencing of Trichophyton rubrum provides insights into population differentiation and drug resistance. Mycopathologia, 2020, 185, 103-112.	1.3	11
27	Molecular and Phenotypic Characterization of Nannizzia (Arthrodermataceae). Mycopathologia, 2020, 185, 9-35.	1.3	14
28	Genome Sequence of the Human Opportunistic Fungus Arthrocladium fulminans (CBS 136243). G3: Genes, Genomes, Genetics, 2020, 10, 1817-1821.	0.8	3
29	<p>Oral Prevalence of Candida Species in Patients Undergoing Systemic Glucocorticoid Therapy and the Antifungal Sensitivity of the Isolates</p> . Infection and Drug Resistance, 2020, Volume 13, 2601-2607.	1.1	10
30	Chromoblastomycosis in an Endemic Area of Brazil: A Clinical-Epidemiological Analysis and a Worldwide Haplotype Network. Journal of Fungi (Basel, Switzerland), 2020, 6, 204.	1.5	11
31	New molecular marker for phylogenetic reconstruction of black yeast-like fungi (Chaetothyriales) with hypothetical EIF2AK2 kinase gene. Fungal Biology, 2020, 124, 1032-1038.	1.1	1
32	Environmental Screening of Fonsecaea Agents of Chromoblastomycosis Using Rolling Circle Amplification. Journal of Fungi (Basel, Switzerland), 2020, 6, 290.	1.5	3
33	Agrobacterium tumefaciens-Mediated Transformation of Fonsecaea monophora and Fonsecaea erecta for Host-Environment Interaction Studies. Journal of Fungi (Basel, Switzerland), 2020, 6, 325.	1.5	2
34	Genomics and Virulence of Fonsecaea pugnacius, Agent of Disseminated Chromoblastomycosis. Frontiers in Genetics, 2020, 11, 822.	1.1	5
35	Onygenalean Fungi as Major Human and Animal Pathogens. Mycopathologia, 2020, 185, 1-8.	1.3	20
36	A re-evaluation of the Chaetothyriales using criteria of comparative biology. Fungal Diversity, 2020, 103, 47-85.	4.7	43

#	Article	IF	CITATIONS
37	No to <i>Neocosmospora</i> : Phylogenomic and Practical Reasons for Continued Inclusion of the Fusarium solani Species Complex in the Genus <i>Fusarium</i> . MSphere, 2020, 5, .	1.3	61
38	Environmental prospecting of black yeast-like agents of human disease using culture-independent methodology. Scientific Reports, 2020, 10, 14229.	1.6	9
39	Comparative Analysis of Clinical and Environmental Strains of Exophiala spinifera by Long-Reads Sequencing and RNAseq Reveal Adaptive Strategies. Frontiers in Microbiology, 2020, 11, 1880.	1.5	6
40	Shed Light in the DaRk LineagES of the Fungal Tree of Life—STRES. Life, 2020, 10, 362.	1.1	16
41	First Case of Rhinocerebral Mucormycosis Caused by Lichtheimia ornata, with a Review of Lichtheimia Infections. Mycopathologia, 2020, 185, 555-567.	1.3	18
42	Selective isolation of agents of chromoblastomycosis from insect-associated environmental sources. Fungal Biology, 2020, 124, 194-204.	1.1	9
43	Comparative Genomic Analysis of Capsule-Producing Black Yeasts Exophiala dermatitidis and Exophiala spinifera, Potential Agents of Disseminated Mycoses. Frontiers in Microbiology, 2020, 11, 586.	1.5	22
44	Intraspecific Diversity and Taxonomy of Emmonsia crescens. Mycopathologia, 2020, 185, 613-627.	1.3	15
45	Transcriptome-wide expression profiling of Sporothrix schenckii yeast and mycelial forms and the establishment of the Sporothrix Genome DataBase. Microbial Genomics, 2020, 6, .	1.0	12
46	Ultrastructure of hyphal cells of the dermatophyte Trichophyton tonsurans. Current Medical Mycology, 2020, 6, 42-46.	0.8	0
47	Prospective application of melanized fungi for the biofiltration of indoor air in closed bioregenerative systems. Journal of Hazardous Materials, 2019, 361, 1-9.	6.5	19
48	New Molecular Markers Distinguishing Fonsecaea Agents of Chromoblastomycosis. Mycopathologia, 2019, 184, 493-504.	1.3	11
49	Changing Concepts and Current Definition of Majocchi's Granuloma. Mycopathologia, 2019, 185, 187-192.	1.3	8
50	A Comparison of Isolation Methods for Black Fungi Degrading Aromatic Toxins. Mycopathologia, 2019, 184, 653-660.	1.3	11
51	Emergomyces: The global rise of new dimorphic fungal pathogens. PLoS Pathogens, 2019, 15, e1007977.	2.1	54
52	Species borderlines in Fusarium exemplified by F. circinatum/F. subglutinans. Fungal Genetics and Biology, 2019, 132, 103262.	0.9	5
53	Species Distinction in the Trichophyton rubrum Complex. Journal of Clinical Microbiology, 2019, 57, .	1.8	35
54	Comparative pathogenicity of opportunistic black yeasts in <i>Aureobasidium</i> . Mycoses, 2019, 62, 803-811.	1.8	16

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55	Virulence and antifungal susceptibility of microsatellite genotypes of <scp><i>Candida albicans</i></scp> from superficial and deep locations. Yeast, 2019, 36, 363-373.	0.8	9
56	Genomic analysis of ant domatia-associated melanized fungi (Chaetothyriales, Ascomycota). Mycological Progress, 2019, 18, 541-552.	0.5	17
57	Peritonitis by Exophiala dermatitidis in a pediatric patient. Medical Mycology Case Reports, 2019, 24, 18-22.	0.7	6
58	Molecular Characterization and Antifungal Susceptibility of Clinical Fusarium Species From Brazil. Frontiers in Microbiology, 2019, 10, 737.	1.5	49
59	Disseminated Mycosis by <i>Arthrocladium</i> <i>fulminans</i> Jeopardizing a Patient with GATA2 Deficiency. Respiration, 2019, 97, 472-475.	1.2	6
60	<p>Multiresistant Fusarium Pathogens on Plants and Humans: Solutions in (from) the Antifungal Pipeline?</p> . Infection and Drug Resistance, 2019, Volume 12, 3727-3737.	1.1	24
61	Epidemiology of <i>Aspergillus</i> species causing keratitis in Mexico. Mycoses, 2019, 62, 144-151.	1.8	25
62	New record of Aureobasidium mangrovei from plant debris in the Sultanate of Oman Czech Mycology, 2019, 71, 219-229.	0.2	3
63	Rapid Identification of Seven Waterborne Exophiala Species by RCA DNA Padlock Probes. Mycopathologia, 2018, 183, 669-677.	1.3	9
64	Black yeasts in the omics era: Achievements and challenges. Medical Mycology, 2018, 56, S32-S41.	0.3	28
65	Phylogeny of dermatophytes with genomic character evaluation of clinically distinct <i>Trichophyton rubrum</i> and <i>T.Âviolaceum</i> . Studies in Mycology, 2018, 89, 153-175.	4.5	50
66	In vitro activities of antifungal drugs against environmental <i>Exophiala</i> isolates and review of the literature. Mycoses, 2018, 61, 561-569.	1.8	12
67	Direct detection of Exophiala and Scedosporium species in sputa of patients with cystic fibrosis. Medical Mycology, 2018, 56, 695-702.	0.3	16
68	Ultra-High-Resolution Mass Spectrometry for Identification of Closely Related Dermatophytes with Different Clinical Predilections. Journal of Clinical Microbiology, 2018, 56, .	1.8	8
69	Genomic Understanding of an Infectious Brain Disease from the Desert. G3: Genes, Genomes, Genetics, 2018, 8, 909-922.	0.8	39
70	Combination of Amphotericin B and Terbinafine against Melanized Fungi Associated with Chromoblastomycosis. Antimicrobial Agents and Chemotherapy, 2018, 62, .	1.4	11
71	New insights in dermatophyte research. Medical Mycology, 2018, 56, S2-S9.	0.3	55
72	Potent Activities of Luliconazole, Lanoconazole, and Eight Comparators against Molecularly Characterized Fusarium Species. Antimicrobial Agents and Chemotherapy, 2018, 62, .	1.4	27

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73	Current antifungal treatment of fusariosis. International Journal of Antimicrobial Agents, 2018, 51, 326-332.	1.1	83
74	Molecular Diagnostics of Arthroconidial Yeasts, Frequent Pulmonary Opportunists. Journal of Clinical Microbiology, 2018, 56, .	1.8	25
75	Antifungal Susceptibility of Emerging Dimorphic Pathogens in the Family Ajellomycetaceae. Antimicrobial Agents and Chemotherapy, 2018, 62, .	1.4	22
76	Fungi between extremotolerance and opportunistic pathogenicity on humans. Fungal Diversity, 2018, 93, 195-213.	4.7	73
77	A Model for Trans-Kingdom Pathogenicity in Fonsecaea Agents of Human Chromoblastomycosis. Frontiers in Microbiology, 2018, 9, 2211.	1.5	20
78	Indoor wet cells as a habitat for melanized fungi, opportunistic pathogens on humans and other vertebrates. Scientific Reports, 2018, 8, 7685.	1.6	15
79	Comparison of the rolling circle amplification and ligase-dependent reaction methods for the identification of opportunistic Exophiala species. Medical Mycology, 2018, 56, 759-769.	0.3	7
80	Global Molecular Diversity of the Halotolerant Fungus Hortaea werneckii. Life, 2018, 8, 31.	1.1	25
81	Phylogeny, ecology and taxonomy of systemic pathogens and their relatives in Ajellomycetaceae (Onygenales): Blastomyces, Emergomyces, Emmonsia, Emmonsiellopsis. Fungal Diversity, 2018, 90, 245-291.	4.7	71
82	Distribution of Pathogens and Outbreak Fungi in the Fungal Kingdom. , 2018, , 3-16.		9
83	Nomenclatural notes on <i>Nadsoniella</i> and the human opportunist black yeast genus <i>Exophiala</i> . Mycoses, 2017, 60, 358-365.	1.8	8
84	Exploring the genomic diversity of black yeasts and relatives (<i>Chaetothyriales</i> , <i>Ascomycota</i>). Studies in Mycology, 2017, 86, 1-28.	4.5	144
85	A novel dimorphic pathogen, <i>Emergomyces orientalis</i> (<i>Onygenales</i>), agent of disseminated infection. Mycoses, 2017, 60, 310-319.	1.8	42
86	Novel taxa of thermally dimorphic systemic pathogens in the <i>Ajellomycetaceae</i> (<i>Onygenales</i>). Mycoses, 2017, 60, 296-309.	1.8	111
87	Diversity of opportunistic black fungi on babassu coconut shells, a rich source of esters and hydrocarbons. Fungal Biology, 2017, 121, 488-500.	1.1	27
88	A phylogenetic perspective on the association between ants (Hymenoptera: Formicidae) and black yeasts (Ascomycota: Chaetothyriales). Proceedings of the Royal Society B: Biological Sciences, 2017, 284, 20162519.	1.2	38
89	First report of urease activity in the novel systemic fungal pathogen Emergomyces africanus: a comparison with the neurotrope Cryptococcus neoformans. FEMS Yeast Research, 2017, 17, .	1.1	11
90	Chromoblastomycosis. Clinical Microbiology Reviews, 2017, 30, 233-276.	5.7	234

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91	Toward a Novel Multilocus Phylogenetic Taxonomy for the Dermatophytes. Mycopathologia, 2017, 182, 5-31.	1.3	447
92	Comparative Evaluation of Etest, EUCAST, and CLSI Methods for Amphotericin B, Voriconazole, and Posaconazole against Clinically Relevant Fusarium Species. Antimicrobial Agents and Chemotherapy, 2017, 61, .	1.4	21
93	Biodiversity and human-pathogenicity of <i> Phialophora verrucosa</i> and relatives in <i> Chaetothyriales</i> . Persoonia: Molecular Phylogeny and Evolution of Fungi, 2017, 38, 1-19.	1.6	26
94	Comparative Genomics of Sibling Species of Fonsecaea Associated with Human Chromoblastomycosis. Frontiers in Microbiology, 2017, 8, 1924.	1.5	31
95	Origin and distribution of Sporothrix globosa causing sapronoses in Asia. Journal of Medical Microbiology, 2017, 66, 560-569.	0.7	62
96	(2465–2466) Proposals to conserve <i>Blastomyces</i> Gilchrist & W. R. Stokes against <i>Blastomyces</i> Costantin & Rolland and <i>Ajellomycetaceae</i> against <i>Paracoccidioidaceae</i> (<i>Ascomycota</i> : <i>Onygenales</i>). Taxon, 2016, 65, 1167-1169.	0.4	6
97	Fusarium: Molecular Diversity and Intrinsic Drug Resistance. PLoS Pathogens, 2016, 12, e1005464.	2.1	314
98	Molecular Epidemiology of Agents of Human Chromoblastomycosis in Brazil with the Description of Two Novel Species. PLoS Neglected Tropical Diseases, 2016, 10, e0005102.	1.3	66
99	Clobal molecular epidemiology and genetic diversity of <i>Fusarium</i> , a significant emerging group of human opportunists from 1958 to 2015. Emerging Microbes and Infections, 2016, 5, 1-11.	3.0	89
100	An updated comprehensive systematic review ofCladophialophora bantianaand analysis of epidemiology, clinical characteristics, and outcome of cerebral cases. Medical Mycology, 2016, 55, myw124.	0.3	45
101	Shared Physiological Traits of Exophiala Species in Cold-Blooded Vertebrates, as Opportunistic Black Yeasts. Mycopathologia, 2016, 181, 353-362.	1.3	1
102	Arthrocladium, an unexpected human opportunist in Trichomeriaceae (Chaetothyriales). Fungal Biology, 2016, 120, 207-218.	1.1	17
103	Draft Genome Sequence of Fonsecaea monophora Strain CBS 269.37, an Agent of Human Chromoblastomycosis. Genome Announcements, 2016, 4, .	0.8	7
104	Draft Genome Sequence of <i>Fonsecaea nubica</i> Strain CBS 269.64, Causative Agent of Human Chromoblastomycosis. Genome Announcements, 2016, 4, .	0.8	6
105	<i>Phaeoacremonium sphinctrophorum</i> as a Novel Agent of Eumycetoma. JAMA Dermatology, 2016, 152, 1063.	2.0	1
106	The role of melanin pathways in extremotolerance and virulence of <i>Fonsecaea</i> revealed by <i>de novo</i> assembly transcriptomics using illumina paired-end sequencing. Studies in Mycology, 2016, 83, 1-18.	4.5	35
107	Barcoding and species recognition of opportunistic pathogens in Ochroconis and Verruconis. Fungal Biology, 2016, 120, 219-230.	1.1	10
108	<i>In vitro</i> combinations of natamycin with voriconazole, itraconazole and micafungin against clinical <i>Fusarium</i> strains causing keratitis: TableÂ1 Journal of Antimicrobial Chemotherapy, 2016, 71, 953-955.	1.3	53

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109	Combination of Amphotericin B and Flucytosine against Neurotropic Species of Melanized Fungi Causing Primary Cerebral Phaeohyphomycosis. Antimicrobial Agents and Chemotherapy, 2016, 60, 2346-2351.	1.4	20
110	Food preparation with mucoralean fungi: A potential biosafety issue?. Fungal Biology, 2016, 120, 393-401.	1.1	19
111	Antifungal Susceptibility and Phylogeny of Opportunistic Members of the Genus <i>Fusarium</i> Causing Human Keratomycosis in South India. Medical Mycology, 2016, 54, 287-294.	0.3	36
112	Bipolaris oryzae, a novel fungal opportunist causing keratitis. Diagnostic Microbiology and Infectious Disease, 2016, 85, 61-65.	0.8	7
113	The â€~species complex' issue in clinically relevant fungi: A case study in Scedosporium apiospermum. Fungal Biology, 2016, 120, 137-146.	1.1	54
114	Extremotolerant rock inhabiting black fungi from Italian monumental sites. Fungal Diversity, 2016, 76, 75-96.	4.7	111
115	DNA barcoding, MALDI-TOF, and AFLP data support Fusarium ficicrescens as a distinct species within the Fusarium fujikuroi species complex. Fungal Biology, 2016, 120, 265-278.	1.1	40
116	Evaluation of two novel barcodes for species recognition of opportunistic pathogens in Fusarium. Fungal Biology, 2016, 120, 231-245.	1.1	48
117	Dishwashers provide a selective extreme environment for human-opportunistic yeast-like fungi. Fungal Diversity, 2016, 76, 1-9.	4.7	52
118	DNA barcoding of fungi causing infections in humans and animals. Fungal Biology, 2016, 120, 125-136.	1.1	67
119	Specific antifungal susceptibility profiles of opportunists in the Fusarium fujikuroi complex. Journal of Antimicrobial Chemotherapy, 2015, 70, 1068-71.	1.3	81
120	<i>In vitro</i> antifungal susceptibility of <i>Trichophyton violaceum</i> isolated from tinea capitis patients. Journal of Antimicrobial Chemotherapy, 2015, 70, 1072-1075.	1.3	10
121	Phylogeography and evolutionary patterns in <1>Sporothrix 1 spanning more than 14 000 human and animal case reports. Persoonia: Molecular Phylogeny and Evolution of Fungi, 2015, 35, 1-20.	1.6	176
122	Evaluation of two molecular techniques for rapid detection of the main dermatophytic agents of tinea capitis. British Journal of Dermatology, 2015, 173, 1494-1500.	1.4	11
123	Rapid Identification of Emerging Human-Pathogenic Sporothrix Species with Rolling Circle Amplification. Frontiers in Microbiology, 2015, 6, 1385.	1.5	38
124	Global Spread of Human Chromoblastomycosis Is Driven by Recombinant Cladophialophora carrionii and Predominantly Clonal Fonsecaea Species. PLoS Neglected Tropical Diseases, 2015, 9, e0004004.	1.3	21
125	One fungus, which genes? Development and assessment of universal primers for potential secondary fungal DNA barcodes. Persoonia: Molecular Phylogeny and Evolution of Fungi, 2015, 35, 242-263.	1.6	416
126	In vitro antifungal susceptibility of coelomycete agents of black grain eumycetoma to eight antifungals. Medical Mycology, 2015, 53, 295-301.	0.3	35

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127	Translation elongation factor $1 \cdot \hat{l} \pm$ gene as a potential taxonomic and identification marker in dermatophytes. Medical Mycology, 2015, 53, 215-224.	0.3	75
128	Black Molds and Melanized Yeasts Pathogenic to Humans. Cold Spring Harbor Perspectives in Medicine, 2015, 5, a019570.	2.9	65
129	Neglected fungal zoonoses: hidden threats to man and animals. Clinical Microbiology and Infection, 2015, 21, 416-425.	2.8	54
130	In VitroInteraction of Currently Used Azoles with Terbinafine against Madurella mycetomatis. Antimicrobial Agents and Chemotherapy, 2015, 59, 1373-1374.	1.4	9
131	MALDI-TOF MS-based identification of black yeasts of the genus Exophiala. Medical Mycology, 2015, 53, 347-352.	0.3	20
132	Epidemiological changes in tinea capitis over the sixty years of economic growth in China. Medical Mycology, 2015, 53, 691-698.	0.3	50
133	Fonsecaea pugnacius, a Novel Agent of Disseminated Chromoblastomycosis. Journal of Clinical Microbiology, 2015, 53, 2674-2685.	1.8	62
134	Phaeohyphomycosis Caused by a Novel Species, Pseudochaetosphaeronema martinelli. Journal of Clinical Microbiology, 2015, 53, 2927-2934.	1.8	24
135	<i>Aspergillus</i> and aspergilloses in wild and domestic animals: a global health concern with parallels to human disease. Medical Mycology, 2015, 53, 765-797.	0.3	172
136	Application of Isothermal Amplification Techniques for Identification of Madurella mycetomatis, the Prevalent Agent of Human Mycetoma. Journal of Clinical Microbiology, 2015, 53, 3280-3285.	1.8	36
137	Draft Genome Sequence of the Ant-Associated Fungus Phialophora attae (CBS 131958). Genome Announcements, 2015, 3, .	0.8	9
138	DNA barcoding of clinically relevant Cunninghamella species. Medical Mycology, 2015, 53, 99-106.	0.3	21
139	Commentaries: Name Changes in Medically Important Fungi and Their Implications for Clinical Practice. Journal of Clinical Microbiology, 2015, 53, 1056-1062.	1.8	65
140	Keratitis by Fusarium temperatum, a novel opportunist. BMC Infectious Diseases, 2014, 14, 588.	1.3	36
141	Black Yeast Diversity on Creosoted Railway Sleepers Changes with Ambient Climatic Conditions. Microbial Ecology, 2014, 68, 699-707.	1.4	38
142	Madurella mycetomatis Is Highly Susceptible to Ravuconazole. PLoS Neglected Tropical Diseases, 2014, 8, e2942.	1.3	43
143	Rapid Identification of Black Grain Eumycetoma Causative Agents Using Rolling Circle Amplification. PLoS Neglected Tropical Diseases, 2014, 8, e3368.	1.3	35
144	Antifungal Susceptibility Patterns of Opportunistic Fungi in the Genera Verruconis and Ochroconis. Antimicrobial Agents and Chemotherapy, 2014, 58, 3285-3292.	1.4	26

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145	Rapid screening for humanâ€pathogenic Mucorales using rolling circle amplification. Mycoses, 2014, 57, 67-72.	1.8	18
146	<i>In Vitro</i> Activities of Eight Antifungal Drugs against 104 Environmental and Clinical Isolates of Aureobasidium pullulans. Antimicrobial Agents and Chemotherapy, 2014, 58, 5629-5631.	1.4	22
147	Environmental siblings of black agents of human chromoblastomycosis. Fungal Diversity, 2014, 65, 47-63.	4.7	56
148	Cyphellophora and its relatives in Phialophora: biodiversity and possible role in human infection. Fungal Diversity, 2014, 65, 17-45.	4.7	62
149	Novel Phialophora species from leaf-cutting ants (tribe Attini). Fungal Diversity, 2014, 65, 65-75.	4.7	39
150	ESCMID and ECMM joint clinical guidelines for the diagnosis and management of systemic phaeohyphomycosis: diseases caused by black fungi. Clinical Microbiology and Infection, 2014, 20, 47-75.	2.8	262
151	Cladophialophora abundans, a novel species of Chaetothyriales isolated from the natural environment. Mycological Progress, 2014, 13, 381-391.	0.5	21
152	Susceptibility and Diversity in the Therapy-Refractory Genus Scedosporium. Antimicrobial Agents and Chemotherapy, 2014, 58, 5877-5885.	1.4	61
153	Black Yeasts and Their Filamentous Relatives: Principles of Pathogenesis and Host Defense. Clinical Microbiology Reviews, 2014, 27, 527-542.	5.7	94
154	Roussoella percutanea, a novel opportunistic pathogen causing subcutaneous mycoses. Medical Mycology, 2014, 52, 689-698.	0.3	26
155	Three Isothermal Amplification Techniques for Rapid Identification of Cladophialophora carrionii, an Agent of Human Chromoblastomycosis. Journal of Clinical Microbiology, 2014, 52, 3531-3535.	1.8	10
156	Implantation phaeohyphomycosis caused by a non-sporulating Chaetomium species. Journal De Mycologie Medicale, 2014, 24, 161-165.	0.7	6
157	Revision of agents of black-grain eumycetoma in the order <i>Pleosporales</i> . Persoonia: Molecular Phylogeny and Evolution of Fungi, 2014, 33, 141-154.	1.6	102
158	Detection and identification of opportunistic Exophiala species using the rolling circle amplification of ribosomal internal transcribed spacers. Journal of Microbiological Methods, 2013, 94, 338-342.	0.7	41
159	Families of Dothideomycetes. Fungal Diversity, 2013, 63, 1-313.	4.7	509
160	Phaeohyphomycoses, Emerging Opportunistic Diseases in Animals. Clinical Microbiology Reviews, 2013, 26, 19-35.	5.7	76
161	<i>In Vitro</i> Antifungal Susceptibility of Cladophialophora carrionii, an Agent of Human Chromoblastomycosis. Antimicrobial Agents and Chemotherapy, 2013, 57, 1974-1977.	1.4	26
162	Chromoblastomycosis caused by Rhinocladiella aquaspersa. Medical Mycology Case Reports, 2013, 2, 148-151.	0.7	25

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163	Black yeast habitat choices and species spectrum onÂhighÂaltitude creosote-treated railway ties. Fungal Biology, 2013, 117, 692-696.	1.1	28
164	Severe Disseminated Phaeohyphomycosis in an Immunocompetent Patient Caused by Veronaea botryosa. Mycopathologia, 2013, 175, 497-503.	1.3	44
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