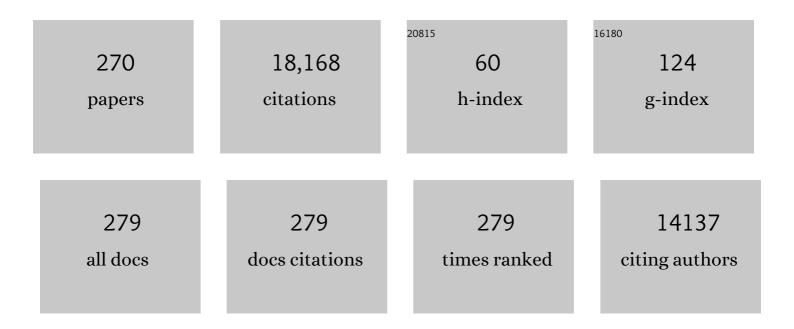
List of Publications by Year in descending order

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FEDDY HACEN

#	Article	IF	CITATIONS
1	Nuclear ribosomal internal transcribed spacer (ITS) region as a universal DNA barcode marker for <i>Fungi</i> . Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 6241-6246.	7.1	4,012
2	Typical freshwater bacteria: an analysis of available 16S rRNA gene sequences from plankton of lakes and rivers. Aquatic Microbial Ecology, 2002, 28, 141-155.	1.8	772
3	A rare genotype of <i>Cryptococcus gattii</i> caused the cryptococcosis outbreak on Vancouver Island (British Columbia, Canada). Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 17258-17263.	7.1	698
4	Recognition of seven species in the Cryptococcus gattii/Cryptococcus neoformans species complex. Fungal Genetics and Biology, 2015, 78, 16-48.	2.1	590
5	First hospital outbreak of the globally emerging Candida auris in a European hospital. Antimicrobial Resistance and Infection Control, 2016, 5, 35.	4.1	535
6	Consensus multi-locus sequence typing scheme for <i>Cryptococcus neoformans</i> and <i>Cryptococcus gattii</i> . Medical Mycology, 2009, 47, 561-570.	0.7	408
7	First report of Candida auris in America: Clinical and microbiological aspects of 18 episodes of candidemia. Journal of Infection, 2016, 73, 369-374.	3.3	340
8	New Clonal Strain of <i>Candida auris</i> , Delhi, India. Emerging Infectious Diseases, 2013, 19, 1670-1673.	4.3	320
9	Multidrug-resistant endemic clonal strain of Candida auris in India. European Journal of Clinical Microbiology and Infectious Diseases, 2014, 33, 919-926.	2.9	303
10	High terbinafine resistance in <i>Trichophyton interdigitale</i> isolates in Delhi, India harbouring mutations in the squalene epoxidase gene. Mycoses, 2018, 61, 477-484.	4.0	237
11	An outbreak due to <i>Candida auris</i> with prolonged colonisation and candidaemia in a tertiary care European hospital. Mycoses, 2018, 61, 498-505.	4.0	236
12	Cryptococcus neoformans-Cryptococcus gattii Species Complex: an International Study of Wild-Type Susceptibility Endpoint Distributions and Epidemiological Cutoff Values for Fluconazole, Itraconazole, Posaconazole, and Voriconazole. Antimicrobial Agents and Chemotherapy, 2012, 56, 5898-5906.	3.2	212
13	The fatal fungal outbreak on Vancouver Island is characterized by enhanced intracellular parasitism driven by mitochondrial regulation. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 12980-12985.	7.1	180
14	Clonal Expansion and Emergence of Environmental Multiple-Triazole-Resistant Aspergillus fumigatus Strains Carrying the TR34/L98H Mutations in the cyp51A Gene in India. PLoS ONE, 2012, 7, e52871.	2.5	180
15	Phylogeography and evolutionary patterns in <i>Sporothrix</i> spanning more than 14 000 human and animal case reports. Persoonia: Molecular Phylogeny and Evolution of Fungi, 2015, 35, 1-20.	4.4	176
16	Genomic Context of Azole Resistance Mutations in Aspergillus fumigatus Determined Using Whole-Genome Sequencing. MBio, 2015, 6, e00536.	4.1	171
17	Six monophyletic lineages identified within Cryptococcus neoformans and Cryptococcus gattii by multi-locus sequence typing. Fungal Genetics and Biology, 2008, 45, 400-421.	2.1	163
18	Proposed nomenclature for Pseudallescheria, Scedosporium and related genera. Fungal Diversity, 2014, 67, 1-10.	12.3	152

#	Article	IF	CITATIONS
19	Diversidad del complejo de especies Cryptococcus neoformans-Cryptococcus gattii. Revista Iberoamericana De Micologia, 2008, 25, S4-S12.	0.9	134
20	Fusarium: more than a node or a foot-shaped basal cell. Studies in Mycology, 2021, 98, 100116.	7.2	134
21	Autochthonous and Dormant <i>Cryptococcus gattii</i> Infections in Europe. Emerging Infectious Diseases, 2012, 18, 1618-1624.	4.3	132
22	Cryptococcus neoformans-Cryptococcus gattii Species Complex: an International Study of Wild-Type Susceptibility Endpoint Distributions and Epidemiological Cutoff Values for Amphotericin B and Flucytosine. Antimicrobial Agents and Chemotherapy, 2012, 56, 3107-3113.	3.2	129
23	Evidence of genotypic diversity among Candida auris isolates by multilocus sequence typing, matrix-assisted laser desorption ionization time-of-flight mass spectrometry and amplified fragment length polymorphism. Clinical Microbiology and Infection, 2016, 22, 277.e1-277.e9.	6.0	127
24	Importance of Resolving Fungal Nomenclature: the Case of Multiple Pathogenic Species in the <i>Cryptococcus</i> Genus. MSphere, 2017, 2, .	2.9	124
25	Paradoxical Immune Responses in Non-HIV Cryptococcal Meningitis. PLoS Pathogens, 2015, 11, e1004884.	4.7	123
26	Unique hybrids between the fungal pathogensCryptococcus neoformansandCryptococcus gattii. FEMS Yeast Research, 2006, 6, 599-607.	2.3	122
27	Ancient Dispersal of the Human Fungal Pathogen Cryptococcus gattii from the Amazon Rainforest. PLoS ONE, 2013, 8, e71148.	2.5	122
28	Multi-azole-resistant Aspergillus fumigatus in the environment in Tanzania. Journal of Antimicrobial Chemotherapy, 2014, 69, 2979-2983.	3.0	122
29	<i>In Vitro</i> Antifungal Susceptibilities and Amplified Fragment Length Polymorphism Genotyping of a Worldwide Collection of 350 Clinical, Veterinary, and Environmental <i>Cryptococcus gattii</i> Isolates. Antimicrobial Agents and Chemotherapy, 2010, 54, 5139-5145.	3.2	121
30	Exploring azole antifungal drug resistance in <i>Aspergillus fumigatus</i> with special reference to resistance mechanisms. Future Microbiology, 2014, 9, 697-711.	2.0	118
31	Azole-resistant Aspergillus fumigatus with the environmental TR46/Y121F/T289A mutation in India. Journal of Antimicrobial Chemotherapy, 2014, 69, 555-557.	3.0	113
32	Candida parapsilosis Resistance to Fluconazole: Molecular Mechanisms and <i>In Vivo</i> Impact in Infected Galleria mellonella Larvae. Antimicrobial Agents and Chemotherapy, 2015, 59, 6581-6587.	3.2	106
33	Tracing Genetic Exchange and Biogeography of <i>Cryptococcus neoformans</i> var. <i>grubii</i> at the Global Population Level. Genetics, 2017, 207, 327-346.	2.9	105
34	A Novel Environmental Azole Resistance Mutation in Aspergillus fumigatus and a Possible Role of Sexual Reproduction in Its Emergence. MBio, 2017, 8, .	4.1	104
35	Global guideline for the diagnosis and management of the endemic mycoses: an initiative of the European Confederation of Medical Mycology in cooperation with the International Society for Human and Animal Mycology. Lancet Infectious Diseases, The, 2021, 21, e364-e374.	9.1	99
36	Environmental study of azoleâ€resistant <i><scp>A</scp>spergillus fumigatus</i> with TR ₃₄ /L98H mutations in the <i>cyp51</i> A gene in <scp>I</scp> ran. Mycoses, 2013, 56, 659-663.	4.0	98

#	Article	IF	CITATIONS
37	Azole-resistant Aspergillus fumigatus harboring TR34/L98H, TR46/Y121F/T289A and TR53 mutations related to flower fields in Colombia. Scientific Reports, 2017, 7, 45631.	3.3	96
38	High prevalence of azole resistance in <i>Aspergillus fumigatus</i> isolates from high-risk patients. Journal of Antimicrobial Chemotherapy, 2015, 70, 2894-2898.	3.0	92
39	AIDS Patient Death Caused by NovelCryptococcus neoformans×C.gattiiHybrid. Emerging Infectious Diseases, 2008, 14, 1105-1108.	4.3	91
40	Prevalence and mechanism of triazole resistance in Aspergillus fumigatus in a referral chest hospital in Delhi, India and an update of the situation in Asia. Frontiers in Microbiology, 2015, 06, 428.	3.5	89
41	Global 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.	6.5	89
42	Molecular characterization of cyanobacterial diversity in a shallow eutrophic lake. Environmental Microbiology, 2005, 7, 365-377.	3.8	87
43	Identification and typing of the emerging pathogen <i>Candida auris</i> by matrixâ€assisted laser desorption ionisation time of flight mass spectrometry. Mycoses, 2016, 59, 535-538.	4.0	86
44	Cryptococcus neoformans Shows a Remarkable Genotypic Diversity in Brazil. Journal of Clinical Microbiology, 2004, 42, 1356-1359.	3.9	83
45	Geographically Structured Populations of Cryptococcus neoformans Variety grubii in Asia Correlate with HIV Status and Show a Clonal Population Structure. PLoS ONE, 2013, 8, e72222.	2.5	83
46	Interaction Between Genetic Background and the Mating-Type Locus in Cryptococcus neoformans Virulence Potential. Genetics, 2005, 171, 975-983.	2.9	82
47	Triazole-resistant Aspergillus fumigatus harbouring G54 mutation: Is it de novo or environmentally acquired?. Journal of Global Antimicrobial Resistance, 2015, 3, 69-74.	2.2	81
48	Beach sand and the potential for infectious disease transmission: observations and recommendations. Journal of the Marine Biological Association of the United Kingdom, 2016, 96, 101-120.	0.8	80
49	Recent trends in molecular diagnostics of yeast infections: from PCR to NGS. FEMS Microbiology Reviews, 2019, 43, 517-547.	8.6	77
50	Passive Surveillance for Azole-Resistant <i>Aspergillus fumigatus</i> , United States, 2011–2013. Emerging Infectious Diseases, 2014, 20, 1498-1503.	4.3	76
51	Low Diversity Cryptococcus neoformans Variety grubii Multilocus Sequence Types from Thailand Are Consistent with an Ancestral African Origin. PLoS Pathogens, 2011, 7, e1001343.	4.7	74
52	Multicenter, International Study of MIC/MEC Distributions for Definition of Epidemiological Cutoff Values for Sporothrix Species Identified by Molecular Methods. Antimicrobial Agents and Chemotherapy, 2017, 61, .	3.2	72
53	Global Population Genetic Analysis of Aspergillus fumigatus. MSphere, 2017, 2, .	2.9	71
54	Nonrandom Distribution of Azole Resistance across the Global Population of Aspergillus fumigatus. MBio, 2019, 10, .	4.1	71

#	Article	IF	CITATIONS
55	Taxonomy and epidemiology of <l>Mucor irregularis</l> , agent of chronic cutaneous mucormycosis. Persoonia: Molecular Phylogeny and Evolution of Fungi, 2013, 30, 48-56.	4.4	69
56	<i>Ceratonia siliqua</i> (carob) trees as natural habitat and source of infection by <i>Cryptococcus gattii</i> in the Mediterranean environment. Medical Mycology, 2012, 50, 67-73.	0.7	67
57	The first cases of <i>Candida auris</i> candidaemia in Oman. Mycoses, 2017, 60, 569-575.	4.0	66
58	Occurrence of triazole-resistant Aspergillus fumigatus with TR34/L98H mutations in outdoor and hospital environment in Kuwait. Environmental Research, 2014, 133, 20-26.	7.5	64
59	Zoonotic transmission of <i>Cryptococcus neoformans</i> from a magpie to an immunocompetent patient. Journal of Internal Medicine, 2005, 257, 385-388.	6.0	63
60	Intercountry Transfer of Triazole-Resistant Aspergillus fumigatus on Plant Bulbs. Clinical Infectious Diseases, 2017, 65, 147-149.	5.8	63
61	Temperate Climate Niche for <i>Cryptococcus gattii</i> in Northern Europe. Emerging Infectious Diseases, 2012, 18, 172-174.	4.3	62
62	Susceptibility and Diversity in the Therapy-Refractory Genus Scedosporium. Antimicrobial Agents and Chemotherapy, 2014, 58, 5877-5885.	3.2	61
63	High prevalence of clinical and environmental triazole-resistant Aspergillus fumigatus in Iran: is it a challenging issue?. Journal of Medical Microbiology, 2016, 65, 468-475.	1.8	60
64	Phylogeny of the industrial relevant, thermophilic genera Myceliophthora and Corynascus. Fungal Diversity, 2012, 52, 197-207.	12.3	59
65	Multicenter Study of Isavuconazole MIC Distributions and Epidemiological Cutoff Values for the Cryptococcus neoformans-Cryptococcus gattii Species Complex Using the CLSI M27-A3 Broth Microdilution Method. Antimicrobial Agents and Chemotherapy, 2015, 59, 666-668.	3.2	58
66	Environmental distribution of <i>Cryptococcus neoformans</i> and <i>C. gattii</i> around the Mediterranean basin. FEMS Yeast Research, 2016, 16, fow045.	2.3	57
67	First Report of Candidemia Clonal Outbreak Caused by Emerging Fluconazole-Resistant Candida parapsilosis Isolates Harboring Y132F and/or Y132F+K143R in Turkey. Antimicrobial Agents and Chemotherapy, 2020, 64, .	3.2	57
68	Attack, Defend and Persist: How the Fungal Pathogen Candida auris was Able to Emerge Globally in Healthcare Environments. Mycopathologia, 2019, 184, 353-365.	3.1	56
69	Emergence of Candida auris in Brazil in a COVID-19 Intensive Care Unit. Journal of Fungi (Basel,) Tj ETQq1 1 0.78	4314 rgBT	- /Qyerlock 1
70	Cryptococcus randhawai sp. nov., a novel anamorphic basidiomycetous yeast isolated from tree trunk hollow of Ficus religiosa (peepal tree) from New Delhi, India. Antonie Van Leeuwenhoek, 2010, 97, 253-259.	1.7	55
71	Concomitant occurrence of itraconazole-resistant and -susceptible strains of Aspergillus fumigatus in routine cultures. Journal of Antimicrobial Chemotherapy, 2015, 70, 412-415.	3.0	55
72	<i>Candida haemulonii</i> species complex: an emerging species in India and its genetic diversity assessed with multilocus sequence and amplified fragment-length polymorphism analyses. Emerging Microbes and Infections, 2016, 5, 1-12.	6.5	55

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73	Prevalence and diversity of filamentous fungi in the airways of cystic fibrosis patients – A Dutch, multicentre study. Journal of Cystic Fibrosis, 2019, 18, 221-226.	0.7	55
74	Identification of Genotypically Diverse Cryptococcus neoformans and Cryptococcus gattii Isolates by Luminex xMAP Technology. Journal of Clinical Microbiology, 2007, 45, 1874-1883.	3.9	54
75	Antifungal susceptibility, genotyping, resistance mechanism, and clinical profile of Candida tropicalis blood isolates. Medical Mycology, 2020, 58, 766-773.	0.7	54
76	Extensive Genetic Diversity within the Dutch Clinical Cryptococcus neoformans Population. Journal of Clinical Microbiology, 2012, 50, 1918-1926.	3.9	53
77	Molecular Epidemiology and In-Vitro Antifungal Susceptibility of Aspergillus terreus Species Complex Isolates in Delhi, India: Evidence of Genetic Diversity by Amplified Fragment Length Polymorphism and Microsatellite Typing. PLoS ONE, 2015, 10, e0118997.	2.5	53
78	Global guidelines and initiatives from the European Confederation of Medical Mycology to improve patient care and research worldwide: New leadership is about working together. Mycoses, 2018, 61, 885-894.	4.0	52
79	Constructing Level-2 Phylogenetic Networks from Triplets. IEEE/ACM Transactions on Computational Biology and Bioinformatics, 2009, 6, 667-681.	3.0	51
80	<i>Cryptococcus gattii</i> Risk for Tourists Visiting Vancouver Island, Canada. Emerging Infectious Diseases, 2007, 13, 178-179.	4.3	49
81	Microsatellite Genotyping Clarified Conspicuous Accumulation of Candida parapsilosis at a Cardiothoracic Surgery Intensive Care Unit. Journal of Clinical Microbiology, 2012, 50, 3422-3426.	3.9	49
82	Invasive Infections Due to <i>Trichosporon</i> : Species Distribution, Genotyping, and Antifungal Susceptibilities from a Multicenter Study in China. Journal of Clinical Microbiology, 2019, 57, .	3.9	49
83	Molecular characterization and <i>in vitro</i> antifungal susceptibility of 80 clinical isolates of mucormycetes in Delhi, India. Mycoses, 2014, 57, 97-107.	4.0	48
84	Home Environment as a Source of Life-Threatening Azole-Resistant <i>Aspergillus fumigatus</i> in Immunocompromised Patients: Table 1 Clinical Infectious Diseases, 2017, 64, 76-78.	5.8	48
85	Prevalence and characterization of azole-resistant Aspergillus fumigatus in patients with cystic fibrosis: a prospective multicentre study in Germany. Journal of Antimicrobial Chemotherapy, 2018, 73, 2047-2053.	3.0	47
86	The global epidemiology of emerging Histoplasma species in recent years. Studies in Mycology, 2020, 97, 100095.	7.2	47
87	Molecular epidemiology and <i>in vitro</i> antifungal susceptibility testing of 108 clinical <i>Cryptococcus neoformans sensu lato</i> and <i>Cryptococcus gattii sensu lato</i> isolates from Denmark. Mycoses, 2016, 59, 576-584.	4.0	46
88	Simple, Low-Cost Detection of Candida parapsilosis Complex Isolates and Molecular Fingerprinting of Candida orthopsilosis Strains in Kuwait by ITS Region Sequencing and Amplified Fragment Length Polymorphism Analysis. PLoS ONE, 2015, 10, e0142880.	2.5	44
89	Fundamental niche prediction of the pathogenic yeasts <i>Cryptococcus neoformans</i> and <i>Cryptococcus gattii</i> in Europe. Environmental Microbiology, 2017, 19, 4318-4325.	3.8	44
90	Evaluation of Molecular Epidemiology, Clinical Characteristics, Antifungal Susceptibility Profiles, and Molecular Mechanisms of Antifungal Resistance of Iranian Candida parapsilosis Species Complex Blood Isolates. Frontiers in Cellular and Infection Microbiology, 2020, 10, 206.	3.9	44

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91	Azole-resistant Aspergillus fumigatus in Denmark: a laboratory-based study on resistance mechanisms and genotypes. Clinical Microbiology and Infection, 2016, 22, 570.e1-570.e9.	6.0	43
92	Routine identification of Nocardia species by MALDI-TOF mass spectrometry. Diagnostic Microbiology and Infectious Disease, 2017, 87, 7-10.	1.8	43
93	YEAST PANEL multiplex PCR for identification of clinically important yeast species: stepwise diagnostic strategy, useful for developing countries. Diagnostic Microbiology and Infectious Disease, 2019, 93, 112-119.	1.8	42
94	Resistance of Asian Cryptococcus neoformans Serotype A Is Confined to Few Microsatellite Genotypes. PLoS ONE, 2012, 7, e32868.	2.5	42
95	Antifungal susceptibility, serotyping, and genotyping of clinical <i>Cryptococcus neoformans</i> isolates collected during 18 years in a single institution in Madrid, Spain. Medical Mycology, 2010, 48, 942-948.	0.7	41
96	Activated dormant <i>Cryptococcus gattii</i> infection in a Dutch tourist who visited Vancouver Island (Canada): a molecular epidemiological approach. Medical Mycology, 2010, 48, 528-531.	0.7	41
97	Comparison of the EUCAST and CLSI Broth Microdilution Methods for Testing Isavuconazole, Posaconazole, and Amphotericin B against Molecularly Identified Mucorales Species. Antimicrobial Agents and Chemotherapy, 2015, 59, 7882-7887.	3.2	41
98	Candida nivariensis Isolated from an Indonesian Human Immunodeficiency Virus-Infected Patient Suffering from Oropharyngeal Candidiasis. Journal of Clinical Microbiology, 2008, 46, 388-391.	3.9	40
99	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.	2.5	40
100	Tuberculosis/cryptococcosis co-infection in China between 1965 and 2016. Emerging Microbes and Infections, 2017, 6, 1-7.	6.5	39
101	Low Level of Antifungal Resistance in Iranian Isolates of Candida glabrata Recovered from Blood Samples in a Multicenter Study from 2015 to 2018 and Potential Prognostic Values of Genotyping and Sequencing of PDR1. Antimicrobial Agents and Chemotherapy, 2019, 63, .	3.2	39
102	The Fungal PCR Initiative's evaluation of in-house and commercial Pneumocystis jirovecii qPCR assays: Toward a standard for a diagnostics assay. Medical Mycology, 2020, 58, 779-788.	0.7	39
103	Itraconazole, Voriconazole, and Posaconazole CLSI MIC Distributions for Wild-Type and Azole-Resistant Aspergillus fumigatus Isolates. Journal of Fungi (Basel, Switzerland), 2018, 4, 103.	3.5	38
104	Novel multiplex real-time quantitative PCR detecting system approach for direct detection of <i>Candida auris</i> and its relatives in spiked serum samples. Future Microbiology, 2019, 14, 33-45.	2.0	38
105	Genotypes and population genetics of cryptococcus neoformans and cryptococcus gattii species complexes in Europe and the mediterranean area. Fungal Genetics and Biology, 2019, 129, 16-29.	2.1	37
106	Species Distinction in the Trichophyton rubrum Complex. Journal of Clinical Microbiology, 2019, 57, .	3.9	35
107	Molecular Identification, Genotypic Diversity, Antifungal Susceptibility, and Clinical Outcomes of Infections Caused by Clinically Underrated Yeasts, Candida orthopsilosis, and Candida metapsilosis: An Iranian Multicenter Study (2014–2019). Frontiers in Cellular and Infection Microbiology, 2019, 9, 264.	3.9	34
108	Successful Allogenic Stem Cell Transplantation in Patients with Inherited CARD9 Deficiency. Journal of Clinical Immunology, 2019, 39, 462-469.	3.8	34

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109	Fungicide-driven alterations in azole-resistant <i>Aspergillus fumigatus</i> are related to vegetable crops in Colombia, South America. Mycologia, 2019, 111, 217-224.	1.9	34
110	Low level of antifungal resistance of <i>Candida glabrata</i> blood isolates in Turkey: Fluconazole minimum inhibitory concentration and <i>FKS</i> mutations can predict therapeutic failure. Mycoses, 2020, 63, 911-920.	4.0	34
111	Molecular epidemiology and antifungal susceptibility of Serbian <i>Cryptococcus neoformans</i> isolates. Mycoses, 2014, 57, 380-387.	4.0	33
112	Microsatellite typing and susceptibilities of serial Cryptococcus neoformansisolates from Cuban patients with recurrent cryptococcal meningitis. BMC Infectious Diseases, 2010, 10, 289.	2.9	32
113	In vitro antimicrobial susceptibility testing of human Brucella melitensis isolates from Qatar between 2014 – 2015. BMC Microbiology, 2015, 15, 121.	3.3	31
114	Cryptococcus tetragattii as a major cause of cryptococcal meningitis among HIV-infected individuals in Harare, Zimbabwe. Journal of Infection, 2016, 72, 745-752.	3.3	31
115	Ecoepidemiology of Cryptococcus gattii in Developing Countries. Journal of Fungi (Basel,) Tj ETQq1 1 0.784314	rgBT_/Ove	rloçk 10 Tf 50
116	Cryptococcal meningitis in systemic lupus erythematosus patients: pooled analysis and systematic review. Emerging Microbes and Infections, 2016, 5, 1-7.	6.5	29
117	Cryptococcosis in patients with diabetes mellitus <scp>II</scp> in mainland China: 1993â€2015. Mycoses, 2017, 60, 706-713.	4.0	29
118	In vitro antifungal activity of amphotericin B and 11 comparators against <i>Aspergillus terreus</i> species complex. Mycoses, 2018, 61, 134-142.	4.0	29
119	Low-Cost Tetraplex PCR for the Global Spreading Multi-Drug Resistant Fungus, Candida auris and Its Phylogenetic Relatives. Frontiers in Microbiology, 2018, 9, 1119.	3.5	29
120	Genetically related micafungin-resistant <i>Candida parapsilosis</i> blood isolates harbouring novel mutation R658G in hotspot 1 of Fks1p: a new challenge?. Journal of Antimicrobial Chemotherapy, 2021, 76, 418-422.	3.0	29
121	Investigation of the basis of virulence in serotype A strains of Cryptococcus neoformans from apparently immunocompetent individuals. Current Genetics, 2004, 46, 92-102.	1.7	28
122	Promiscuous mitochondria in <i>Cryptococcus gattii</i> . FEMS Yeast Research, 2009, 9, 489-503.	2.3	28
123	Internal validation of <scp>CPS</scp> ^{â,,¢} <scp>MONODOSE</scp> CanAur dtecâ€ <scp>qPCR</scp> kit following the <scp>UNE</scp> / <scp>EN ISO</scp> / <scp>IEC</scp> 17025:2005 for detection of the emerging yeast <i>Candida auris</i> . Mycoses, 2018, 61, 877-884.	4.0	28
124	Comparison of 21-Plex PCR and API 20C AUX, MALDI-TOF MS, and rDNA Sequencing for a Wide Range of Clinically Isolated Yeast Species: Improved Identification by Combining 21-Plex PCR and API 20C AUX as an Alternative Strategy for Developing Countries. Frontiers in Cellular and Infection Microbiology, 2019, 9, 21.	3.9	28
125	Performance of Two Novel Chromogenic Media for the Identification of Multidrug-Resistant Candida auris Compared with Other Commercially Available Formulations. Journal of Clinical Microbiology, 2021, 59, .	3.9	28
126	Cryptococcus neoformans population diversity and clinical outcomes of HIV-associated cryptococcal meningitis patients in Zimbabwe. Journal of Medical Microbiology, 2016, 65, 1281-1288.	1.8	28

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127	Interlaboratory evaluation of Mucorales PCR assays for testing serum specimens: A study by the fungal PCR Initiative and the Modimucor study group. Medical Mycology, 2021, 59, 126-138.	0.7	27
128	Triazole phenotypes and genotypic characterization of clinical <i>Aspergillus fumigatus</i> isolates in China. Emerging Microbes and Infections, 2017, 6, 1-6.	6.5	26
129	Outbreak of Fusarium oxysporum infections in children with cancer: an experience with 7 episodes of catheter-related fungemia. Antimicrobial Resistance and Infection Control, 2017, 6, 93.	4.1	26
130	The Search for the Natural Habitat of Cryptococcus gattii. Mycopathologia, 2010, 170, 209-211.	3.1	25
131	Molecular Diagnostics of Arthroconidial Yeasts, Frequent Pulmonary Opportunists. Journal of Clinical Microbiology, 2018, 56, .	3.9	25
132	Global Molecular Diversity of the Halotolerant Fungus Hortaea werneckii. Life, 2018, 8, 31.	2.4	25
133	In vitro antifungal susceptibilities and molecular typing of sequentially isolated clinical Cryptococcus neoformans strains from Croatia. Journal of Medical Microbiology, 2011, 60, 1487-1495.	1.8	24
134	Molecular characterisation and antifungal susceptibility of clinical Cryptococcus deuterogattii (AFLP6/VGII) isolates from Southern Brazil. European Journal of Clinical Microbiology and Infectious Diseases, 2016, 35, 1803-1810.	2.9	24
135	Determining the analytical specificity of PCR-based assays for the diagnosis of IA: What is <i>Aspergillus</i> ?. Medical Mycology, 2017, 55, myw093.	0.7	24
136	Growth and Mating of Cryptococcus neoformans var. grubii on Woody Debris. Microbial Ecology, 2009, 57, 757-765.	2.8	23
137	Pitfalls in Serological Diagnosis ofCryptococcus gattiiInfections. Medical Mycology, 2015, 53, 874-879.	0.7	23
138	Geographically predominant genotypes of Aspergillus terreus species complex in Austria: s microsatellite typing study. Clinical Microbiology and Infection, 2016, 22, 270-276.	6.0	23
139	High-resolution fingerprinting of Candida parapsilosis isolates suggests persistence and transmission of infections among neonatal intensive care unit patients in Kuwait. Scientific Reports, 2019, 9, 1340.	3.3	23
140	Meningitis caused by <i>Filobasidium uniguttulatum</i> : case report and overview of the literature. Mycoses, 2012, 55, 105-109.	4.0	22
141	First environmental isolation of <i>Cryptococcus gattii</i> , genotype AFLP5, from India and a global review. Mycoses, 2013, 56, 222-228.	4.0	22
142	Coxiella burnetii Genotypes in Iberian Wildlife. Microbial Ecology, 2016, 72, 890-897.	2.8	22
143	Cenome-wide mapping using new AFLP markers to explore intraspecific variation among pathogenic Sporothrix species. PLoS Neglected Tropical Diseases, 2020, 14, e0008330.	3.0	22
144	Global Spread of Human Chromoblastomycosis Is Driven by Recombinant Cladophialophora carrionii and Predominantly Clonal Fonsecaea Species. PLoS Neglected Tropical Diseases, 2015, 9, e0004004.	3.0	21

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145	Environmental distribution of <i>Cryptococcus</i> species and some other yeastâ€kke fungi in India. Mycoses, 2018, 61, 305-313.	4.0	21
146	Eighty Years of Mycopathologia: A Retrospective Analysis of Progress Made in Understanding Human and Animal Fungal Pathogens. Mycopathologia, 2018, 183, 859-877.	3.1	21
147	Evaluation of Microsatellite Typing, ITS Sequencing, AFLP Fingerprinting, MALDI-TOF MS, and Fourier-Transform Infrared Spectroscopy Analysis of Candida auris. Journal of Fungi (Basel,) Tj ETQq1 1 0.78431	4 rg₿₮ /Ov	verlæak 10 Tf
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