Carolina Firacative

List of Publications by Year
in descending order

59 papers	1,412 citations	411340 20 h-index	388640 36 g-index
60 all docs	60 docs citations	60 times ranked	1643 citing authors

#	Article	IF	Citations
1	Recent Advances in Cryptococcus and Cryptococcosis. Microorganisms, 2022, 10, 13.	1.6	10
2	Molecular type distribution and fluconazole susceptibility of clinical Cryptococcus gattii isolates from South African laboratory-based surveillance, 2005â€“2013. PLoS Neglected Tropical Diseases, 2022, 16, e0010448.	1.3	1
3	A case-series of bloodstream infections caused by theMeyerozyma guilliermondiispecies complex at a reference center of oncology in Brazil. Medical Mycology, 2021, 59, 235-243.	0.3	13
4	Genotype, Antifungal Susceptibility, and Virulence of Clinical South African Cryptococcus neoformans Strains from National Surveillance, 2005â€“2009. Journal of Fungi (Basel, Switzerland), 2021, 7, 338.	1.5	5
5	Cryptococcus neoformans and Cryptococcus gattii Species Complexes in Latin America: A Map of Molecular Types, Genotypic Diversity, and Antifungal Susceptibility as Reported by the Latin American Cryptococcal Study Group. Journal of Fungi (Basel, Switzerland), 2021, 7, 282.	1.5	20
6	Antifungal susceptibility of clinical Cryptococcus gattii isolates from Colombia varies among molecular types. Medical Mycology, 2021, 59, 1122-1125.	0.3	6
7	Identification of Disease-Associated Cryptococcal Proteins Reactive With Serum IgG From Cryptococcal Meningitis Patients. Frontiers in Immunology, 2021, 12, 709695.	2.2	8
8	Cryptococcus neoformans VNII as the Main Cause of Cryptococcosis in Domestic Cats from Rio de Janeiro, Brazil. Journal of Fungi (Basel, Switzerland), 2021, 7, 980.	1.5	4
9	Multilocus Sequence Typing Reveals Extensive Genetic Diversity of the Emerging Fungal Pathogen Scedosporium aurantiacum. Frontiers in Cellular and Infection Microbiology, 2021, 11, 761596.	1.8	4
10	Clinical and Epidemiological Profile of Patients with Invasive Aspergillosis from a Fourth Level Hospital in Bogota, Colombia: A Retrospective Study. Journal of Fungi (Basel, Switzerland), 2021, 7, 1092.	1.5	3
11	Molecular identification and antifungal susceptibility testing of Pucciniomycotina red yeast clinical isolates from Rio de Janeiro, Brazil. Brazilian Journal of Microbiology, 2020, 51, 95-98.	0.8	2
12	Molecular Epidemiology Reveals Low Genetic Diversity among Cryptococcus neoformans Isolates from People Living with HIV in Lima, Peru, during the Pre-HAART Era. Pathogens, 2020, 9, 665.	1.2	7
13	Rearing and Maintenance of Galleria mellonella and Its Application to Study Fungal Virulence. Journal of Fungi (Basel, Switzerland), 2020, 6, 130.	1.5	32
14	Cryptococcosis in Hematopoietic Stem Cell Transplant Recipients: A Rare Presentation Warranting Recognition. Canadian Journal of Infectious Diseases and Medical Microbiology, 2020, 2020, 1-8.	0.7	6
15	Indoor Dust as a Source of Virulent Strains of the Agents of Cryptococcosis in the Rio Negro Micro-Region of the Brazilian Amazon. Microorganisms, 2020, 8, 682.	1.6	8
16	A screening of the MMV Pathogen BoxÂ® reveals new potential antifungal drugs against the etiologic agents of chromoblastomycosis. PLoS ONE, 2020, 15, e0229630.	1.1	18
17	Invasive fungal disease in humans: are we aware of the real impact?. Memorias Do Instituto Oswaldo Cruz, 2020, 115, e200430.	0.8	96
18	Comparative Analysis of Putative Virulence-Associated Factors of Microsporum canis Isolates from Human and Animal Patients. Mycopathologia, 2020, 185, 665-673.	1.3	5

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19	Fatal fungaemia due to Cryptococcus albidus in an elderly diabetic woman presenting with pleural effusion. Revista Do Instituto De Medicina Tropical De Sao Paulo, 2020, 62, e34.	0.5	3
20	Title is missing!. , 2020, 15, e0229630.		0
21	Title is missing!. , 2020, 15, e0229630.		0
22	Title is missing!. , 2020, 15, e0229630.		0
23	Title is missing!. , 2020, 15, e0229630.		0
24	Title is missing!. , 2020, 15, e0229630.		0
25	Title is missing!. , 2020, 15, e0229630.		0
26	Comparative antifungal susceptibility analyses of Cryptococcus neoformans VNI and Cryptococcus gattii VGII from the Brazilian Amazon Region by the Etest, Vitek 2, and the Clinical and Laboratory Standards Institute broth microdilution methods. Medical Mycology, 2019, 57, 864-873.	0.3	10
27	Clonal Dispersal of Cryptococcus gattii VGII in an Endemic Region of Cryptococcosis in Colombia. Journal of Fungi (Basel, Switzerland), 2019, 5, 32.	1.5	10
28	Cryptococcosis due to Cryptococcus gattii VGII in southeast Brazil: The One Health approach revealing a possible role for domestic cats. Medical Mycology Case Reports, 2019, 24, 61-64.	0.7	9
29	Identification of T helper (Th)1- and Th2-associated antigens of Cryptococcus neoformans in a murine model of pulmonary infection. Scientific Reports, 2018, 8, 2681.	1.6	73
30	Molecular identification and antifungal susceptibility profiles of clinical strains of Fonsecaea spp. isolated from patients with chromoblastomycosis in Rio de Janeiro, Brazil. PLoS Neglected Tropical Diseases, 2018, 12, e0006675.	1.3	23
31	The status of cryptococcosis in Latin America. Memorias Do Instituto Oswaldo Cruz, 2018, 113, e170554.	0.8	66
32	Hepatic Disease with Portal Hypertension and Acute Juvenile Paracoccidioidomycosis: A Report of Two Cases and Literature Review. Mycopathologia, 2017, 182, 915-919.	1.3	7
33	Tinea Capitis by Microsporum audouinii: Case Reports and Review of Published Global Literature 2000â€“2016. Mycopathologia, 2017, 182, 1053-1060.	1.3	18
34	Advances in the understanding of the Cryptococcus neoformans and C. gattii species complexes and cryptococcosis. Microbiology Australia, 2017, 38, 106.	0.1	1
35	MLST and Whole-Genome-Based Population Analysis of Cryptococcus gattii VGIII Links Clinical, Veterinary and Environmental Strains, and Reveals Divergent Serotype Specific Sub-populations and Distant Ancestors. PLoS Neglected Tropical Diseases, 2016, 10, e0004861.	1.3	49
36	Population Genetic Analysis Reveals a High Genetic Diversity in the Brazilian Cryptococcus gattii VGII Population and Shifts the Global Origin from the Amazon Rainforest to the Semi-arid Desert in the Northeast of Brazil. PLoS Neglected Tropical Diseases, 2016, 10, e0004885.	1.3	52

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37	Pathogenic diversity amongst serotype C VGIII and VGIV Cryptococcus gattii isolates. Scientific Reports, 2015, 5, 11717.	1.6	15
38	Environmental Isolation of Cryptococcus gattii VGII from Indoor Dust from Typical Wooden Houses in the Deep Amazonas of the Rio Negro Basin. PLoS ONE, 2015, 10, e0115866.	1.1	42
39	Australia in the global picture of the molecular epidemiology of Cryptococcus gattii molecular type VGII. Microbiology Australia, 2015, 36, 67.	0.1	3
40	Fatal Case of Polymicrobial Meningitis Caused by Cryptococcus liquefaciens and Mycobacterium tuberculosis Complex in a Human Immunodeficiency Virus-Infected Patient. Journal of Clinical Microbiology, 2015, 53, 2753-2755.	1.8	16
41	Galleria mellonella Model Identifies Highly Virulent Strains among All Major Molecular Types of Cryptococcus gattii. PLoS ONE, 2014, 9, e105076.	1.1	56
42	Cryptococcus gattii in North American Pacific Northwest: Whole-Population Genome Analysis Provides Insights into Species Evolution and Dispersal. MBio, 2014, 5, e01464-14.	1.8	126
43	Retrospective Study of the Epidemiology and Clinical Manifestations of Cryptococcus gattii Infections in Colombia from 1997â€“2011. PLoS Neglected Tropical Diseases, 2014, 8, e3272.	1.3	51
44	Phenotypic Differences of Cryptococcus Molecular Types and Their Implications for Virulence in a Drosophila Model of Infection. Infection and Immunity, 2014, 82, 3058-3065.	1.0	33
45	MALDI-TOF MS for the identification of veterinary non-C. neoformans-C. gattii Cryptococcus spp. isolates from Italy. Medical Mycology, 2014, 52, 659-666.	0.3	4
46	Multilocus sequence typing (MLST) and M13 PCR fingerprinting revealed heterogeneity amongstCryptococcusspecies obtained from Italian veterinary isolates. FEMS Yeast Research, 2014, 14, 897-909.	1.1	36
47	Antifungal Drug Susceptibility and Phylogenetic Diversity among Cryptococcus Isolates from Dogs and Cats in North America. Journal of Clinical Microbiology, 2014, 52, 2061-2070.	1.8	40
48	Hospital-acquired Pneumocystis pneumonia: a renewed concern?. Microbiology Australia, 2014, 35, 57.	0.1	1
49	Identification of the Major Molecular Types of Cryptococcus neoformans and C. gattii by Hyperbranched Rolling Circle Amplification. PLoS ONE, 2014, 9, e94648.	1.1	39
50	Molecular Epidemiology Reveals Genetic Diversity amongst Isolates of the Cryptococcus neoformans/C. gattii Species Complex in Thailand. PLoS Neglected Tropical Diseases, 2013, 7, e2297.	1.3	54
51	Molecular Epidemiology Linking Multihospital Clusters of Opportunistic Pneumocystis jirovecii Pneumonia. Clinical Infectious Diseases, 2013, 57, 1058-1059.	2.9	17
52	MALDI-TOF MS Enables the Rapid Identification of the Major Molecular Types within the Cryptococcus neoformans/C. gattii Species Complex. PLoS ONE, 2012, 7, e37566.	1.1	113
53	Primer aislamiento ambiental de Cryptococcus gattii de serotipo B, en CÃºcuta, Colombia. Biomedica, 2011, 31, 118.	0.3	22
54	Nosocomial Pneumocystis jirovecii Pneumonia: Lessons From a Cluster in Kidney Transplant Recipients. Transplantation, 2011, 92, 1327-1334.	0.5	82

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55	Fatal Disseminated Cryptococcus gattii Infection in New Mexico. PLoS ONE, 2011, 6, e28625.	1.1	38
56	First environmental isolation of Cryptococcus gattii serotype B, from CÃºcuta, Colombia. Biomedica, 2011, 31, 118-23.	0.3	19
57	Isolation ofCryptococcus gattiimolecular type VGIII, fromCorymbia ficifoliadetritus in Colombia. Medical Mycology, 2010, 48, 675-678.	0.3	28
58	Circulation of Streptococcus pneumoniae clone Colombia5 ST289 in nine Latin American countries. Revista Panamericana De Salud Publica/Pan American Journal of Public Health, 2009, 25, 337-43.	0.6	7
59	CaracterizaciÃ³n molecular de aislamientos invasores colombianos de Streptococcus pneumoniae serotipo 5 recuperados entre 1994 y 2004.. Biomedica, 2006, 26, 295.	0.3	1