Massimo Cogliati

List of Publications by Year
in descending order

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49 papers	2,083 citations	331670 21 h-index	243625 44 g-index
51 all docs	51 docs citations	51 times ranked	1701 citing authors

#	Article	IF	Citations
1	Consensus multi-locus sequence typing scheme forCryptococcus neoformansandCryptococcus gattii. Medical Mycology, 2009, 47, 561-570.	0.7	408
2	The Case for Adopting the â€œSpecies Complexâ€•Nomenclature for the Etiologic Agents of Cryptococcosis. MSphere, 2017, 2, .	2.9	274
3	Global Molecular Epidemiology ofCryptococcus neoformansandCryptococcus gattii: An Atlas of the Molecular Types. Scientifica, 2013, 2013, 1-23.	1.7	226
4	Molecular analysis of 311Cryptococcus neoformansisolates from a 30-month ECMM survey of cryptococcosis in Europe. FEMS Yeast Research, 2006, 6, 614-619.	2.3	134
5	Autochthonous and DormantCryptococcus gattiiInfections in Europe. Emerging Infectious Diseases, 2012, 18, 1618-1624.	4.3	132
6	Matrix-Assisted Laser Desorption Ionizationâ€“Time of Flight Mass Spectrometry-Based Method for Discrimination between Molecular Types of Cryptococcus neoformans and Cryptococcus gattii. Journal of Clinical Microbiology, 2012, 50, 2472-2476.	3.9	87
7	Origin of Cryptococcus neoformans var. neoformans Diploid Strains. Journal of Clinical Microbiology, 2001, 39, 3889-3894.	3.9	73
8	MLST-Based Population Genetic Analysis in a Global Context Reveals Clonality amongst Cryptococcus neoformans var. grubii VNI Isolates from HIV Patients in Southeastern Brazil. PLoS Neglected Tropical Diseases, 2017, 11, e0005223.	3.0	59
9	Environmental distribution ofCryptococcus neoformansandC. gattiiaround the Mediterranean basin. FEMS Yeast Research, 2016, 16, fow045.	2.3	57
10	Fundamental niche prediction of the pathogenic yeasts Cryptococcus neoformans and Cryptococcus gattii in Europe. Environmental Microbiology, 2017, 19, 4318-4325.	3.8	44
11	Cryptococcus neoformanspopulation includes hybrid strains homozygous at mating-type locus. FEMS Yeast Research, 2006, 6, 608-613.	2.3	39
12	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
13	Multilocus sequence typing (MLST) and M13 PCR fingerprinting revealed heterogeneity amongstCryptococcusspecies obtained from Italian veterinary isolates. FEMS Yeast Research, 2014, 14, 897-909.	2.3	36
14	Multilocus sequence typing analysis reveals that Cryptococcus neoformans var. neoformans is a recombinant population. Fungal Genetics and Biology, 2016, 87, 22-29.	2.1	34
15	Molecular epidemiology of Italian clinicalCryptococcus neoformansvar.grubiiisolates. Medical Mycology, 2013, 51, 499-506.	0.7	33
16	Azole Resistance in Aspergillus fumigatus Clinical Isolates from an Italian Culture Collection. Antimicrobial Agents and Chemotherapy, 2016, 60, 682-685.	3.2	32
17	Isolation, Identification and Molecular Typing of Cryptococcus neoformans from Pigeon Droppings and Other Environmental Sources in Tripoli, Libya. Mycopathologia, 2016, 181, 603-608.	3.1	29
18	Cryptococcus gattii serotypeâ€€ strains isolated in Bangalore, Karnataka, India. Mycoses, 2012, 55, 262-268.	4.0	25

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19	Four-Year Persistence of a Single Candida albicans Genotype Causing Bloodstream Infections in a Surgical Ward Proven by Multilocus Sequence Typing. Journal of Clinical Microbiology, 2006, 44, 218-221.	3.9	24
20	Determining the analytical specificity of PCR-based assays for the diagnosis of IA: What isAspergillus?. Medical Mycology, 2017, 55, myw093.	0.7	24
21	Molecular characterization of C ryptococcus neoformans and C ryptococcus gattii from environmental sources and genetic comparison with clinical isolates in Apulia, Italy. Environmental Research, 2018, 160, 347-352.	7.5	24
22	Cryptococcus gattii infection in an immunocompetent host in Greece. Medical Mycology Case Reports, 2020, 27, 1-3.	1.3	22
23	Heterozygosis and Pathogenicity of Cryptococcus neoformans AD-Hybrid Isolates. Mycopathologia, 2012, 173, 347-357.	3.1	20
24	Global warming impact on the expansion of fundamental niche of Cryptococcus gattii VGI in Europe. Environmental Microbiology Reports, 2021, 13, 375-383.	2.4	19
25	Comparison of Three Methods for Testing Azole Susceptibilities of Candida albicans Strains Isolated Sequentially from Oral Cavities of AIDS Patients. Journal of Clinical Microbiology, 1998, 36, 1578-1583.	3.9	17
26	Molecular characterization of environmental Cryptococcus neoformans VNII isolates in Jos, Plateau State, Nigeria. Journal De Mycologie Medicale, 2016, 26, 306-311.	1.5	16
27	Genetic Factors and Genotype-Environment Interactions Contribute to Variation in Melanin Production in the Fungal Pathogen Cryptococcus neoformans. Scientific Reports, 2018, 8, 9824.	3.3	16
28	Epidemiological trends of cryptococcosis in Italy: Molecular typing and susceptibility pattern of Cryptococcus neoformans isolates collected during a 20-year period. Medical Mycology, 2018, 56, 963-971.	0.7	12
29	Identification and Characterization of VNI/VNII and Novel VNII/VNIV Hybrids and Impact of Hybridization on Virulence and Antifungal Susceptibility Within the C. neoformans/C. gattii Species Complex. PLoS ONE, 2016, 11, e0163955.	2.5	12
30	Spatial Quantification of the Population Exposed to Cryptococcus neoformans and Cryptococcus gattii Species Complexes in Europe: Estimating the Immunocompetent and HIV/AIDS Patients Under Risk. Risk Analysis, 2020, 40, 524-533.	2.7	10
31	Epidemiological characteristics of cryptococcal meningoencephalitis associated with Cryptococcus neoformans var. grubii from HIV-infected patients in Madagascar: A cross-sectional study. PLoS Neglected Tropical Diseases, 2020, 14, e0007984.	3.0	10
32	First report of two cases of cryptococcosis in Tripoli, Libya, infected with Cryptococcus neoformans isolates present in the urban area. Journal De Mycologie Medicale, 2017, 27, 421-424.	1.5	9
33	Azole resistance in Aspergillus isolates by different types of patients and correlation with environment â€•An Italian prospective multicentre study (ARiA study). Mycoses, 2021, 64, 528-536.	4.0	9
34	The need for environmental surveillance to understand the ecology, epidemiology and impact of Cryptococcus infection in Africa. FEMS Microbiology Ecology, 2021, 97, .	2.7	9
35	Cryptococcus gattii sero-mating type allelic pattern determined by multiplex PCR. Clinical Microbiology and Infection, 2015, 21, 190.e1-190.e4.	6.0	7
36	Expansion of the Emerging Fungal Pathogen Cryptococcus bacillisporus Into America: Linking Phylogenetic Origin, Geographical Spread and Population Under Exposure Risk. Frontiers in Microbiology, 2020, 11, 2117.	3.5	7

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37	Cryptococcus neoformans Typing by PCR Fingerprinting Using (GACA) ₄ Primers Based on C. neoformans Genome Project Data. Journal of Clinical Microbiology, 2007, 45, 3427-3430.	3.9	6
38	Electrophoretic karyotyping of Cryptococcus neoformans ADâ€hybrid strains. Mycoses, 2009, 52, 16-23.	4.0	5
39	Molecular-Type Specific Multiplex PCR produces a distinct VNII PCR pattern amongCryptococcus neoformansspecies complex. Medical Mycology, 2019, 57, 384-386.	0.7	5
40	Cryptococcus neoformans and Cryptococcus gattii Species Complex Isolates on the Slopes of Mount Etna, SICILY, Italy. Frontiers in Microbiology, 2019, 10, 2390.	3.5	4
41	First Isolation, Antifungal Susceptibility, and Molecular Characterization of Cryptococcus neoformans from the Environment in Croatia. Journal of Fungi (Basel, Switzerland), 2019, 5, 99.	3.5	4
42	Cryptococcus neoformans species complex isolates living in a tree micro-ecosystem. Fungal Ecology, 2020, 44, 100889.	1.6	4
43	New multilocus sequence typing primers to enable genotyping of AD hybrids within the Cryptococcus neoformans species complex. Medical Mycology, 2020, 58, 1005-1009.	0.7	4
44	A case of Histoplasma capsulatum endophthalmitis diagnosed in Italy. Travel Medicine and Infectious Disease, 2013, 11, 256-258.	3.0	3
45	Comment on: T2Candida MR as a predictor of outcome in patients with suspected invasive candidiasis starting empirical antifungal treatment: a prospective pilot study. Journal of Antimicrobial Chemotherapy, 2019, 74, 532-533.	3.0	3
46	Trichophyton tonsurans in Chile: Genotyping in search of an origin. Medical Mycology, 2022, 60, .	0.7	3
47	First Autochthonous Case of Cryptococcal Meningitis in an Immunocompetent Host Due to Cryptococcus gattii VGIin Northern Italy. SN Comprehensive Clinical Medicine, 2020, 2, 237-241.	0.6	2
48	Hybridization and Its Importance in the Cryptococcus Species Complex. , 0, , 359-370.		2
49	Multi-locus sequence typing reveals genotypic similarity in Nigerian Cryptococcus neoformans AFLP1/VNI of environmental and clinical origin. Journal of Medical Microbiology, 2021, 70, .	1.8	2