## Marie Desnos-Ollivier

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6767833/publications.pdf

Version: 2024-02-01

55 papers 3,008 citations

30 h-index 53 g-index

59 all docs 59 docs citations

59 times ranked

3688 citing authors

#	Article	IF	CITATIONS
1	Bloodstream Infections Caused by <i>Magnusiomyces capitatus</i> and <i>Magnusiomyces clavatus</i> : Epidemiological, Clinical, and Microbiological Features of Two Emerging Yeast Species. Antimicrobial Agents and Chemotherapy, 2022, 66, AAC0183421.	3.2	10
2	Echinocandins Susceptibility Patterns of 2,787 Yeast Isolates: Importance of the Thresholds for the Detection of FKS Mutations. Antimicrobial Agents and Chemotherapy, 2022, 66, e0172521.	3.2	6
3	Active Surveillance Program to Increase Awareness on Invasive Fungal Diseases: the French RESSIF Network (2012 to 2018). MBio, 2022, 13, e0092022.	4.1	26
4	Central nervous system candidiasis beyond neonates: Lessons from a nationwide study. Medical Mycology, 2021, 59, 266-277.	0.7	15
5	A Short-Tandem-Repeat Assay ( Mmy STR) for Studying Genetic Variation in Madurella mycetomatis. Journal of Clinical Microbiology, 2021, 59, .	3.9	6
6	Case Report: Emergence of Candida auris in the Indian Ocean Region. American Journal of Tropical Medicine and Hygiene, 2021, 104, 739-743.	1.4	2
7	Cryptococcus gattii in Patients with Lymphoid Neoplasms: An Illustration of Evolutive Host–Fungus Interactions. Journal of Fungi (Basel, Switzerland), 2021, 7, 212.	3.5	4
8	No Impact of Fluconazole to Echinocandins Replacement as First-Line Therapy on the Epidemiology of Yeast Fungemia (Hospital-Driven Active Surveillance, 2004–2017, Paris, France). Frontiers in Medicine, 2021, 8, 641965.	2.6	8
9	Azole Susceptibility Profiles of More than 9,000 Clinical Yeast Isolates Belonging to 40 Common and Rare Species. Antimicrobial Agents and Chemotherapy, 2021, 65, .	3.2	17
10	Kazachstania slooffiae: An unexpected journey to a human pleural sample. Journal De Mycologie Medicale, 2021, 31, 101109.	1.5	7
11	Earliest case of Candida auris infection imported in 2007 in Europe from India prior to the 2009 description in Japan. Journal De Mycologie Medicale, 2021, 31, 101139.	1.5	16
12	Cryptococcus gattii Species Complex as an Opportunistic Pathogen: Underlying Medical Conditions Associated with the Infection. MBio, 2021, 12, e0270821.	4.1	25
13	The genus Madurella: Molecular identification and epidemiology in Sudan. PLoS Neglected Tropical Diseases, 2020, 14, e0008420.	3.0	8
14	Comparison of MultiLocus Sequence Typing (MLST) and Microsatellite Length Polymorphism (MLP) for Pneumocystis jirovecii genotyping. Computational and Structural Biotechnology Journal, 2020, 18, 2890-2896.	4.1	5
15	<i>Saprochaete clavata</i> Outbreak Infecting Cancer Center through Dishwasher. Emerging Infectious Diseases, 2020, 26, 2031-2038.	4.3	17
16	Epidemiological investigation for grouped cases of Trichosporon asahii using whole genome and IGS1 sequencing. Mycoses, 2020, 63, 942-951.	4.0	5
17	Madurella real-time PCR, a novel approach for eumycetoma diagnosis. PLoS Neglected Tropical Diseases, 2020, 14, e0007845.	3.0	9
18	<i>Yarrowia lipolytica</i> causes sporadic cases and local outbreaks of infections and colonisation. Mycoses, 2020, 63, 737-745.	4.0	12

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19	Tracing the Evolutionary History and Global Expansion of Candida auris Using Population Genomic Analyses. MBio, 2020, $11$ , .	4.1	224
20	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
21	Database establishment for the secondary fungal DNA barcode (i>translational elongation factor $1\hat{1}\pm\langle i \rangle$ ((i>TEF1 $\hat{1}\pm\langle i \rangle$ ). Genome, 2019, 62, 160-169.	2.0	41
22	Population Structure of Candida parapsilosis: No Genetic Difference Between French and Uruguayan Isolates Using Microsatellite Length Polymorphism. Mycopathologia, 2018, 183, 381-390.	3.1	8
23	Investigating Clinical Issues by Genotyping of Medically Important Fungi: Why and How?. Clinical Microbiology Reviews, 2017, 30, 671-707.	13.6	65
24	Rearranged Biosynthetic Gene Cluster and Synthesis of Hassallidin E in <i>Planktothrix serta</i> PCC 8927. ACS Chemical Biology, 2017, 12, 1796-1804.	3.4	25
25	Predisposing factors and outcome of uncommon yeast species-related fungaemia based on an exhaustive surveillance programme (2002–14). Journal of Antimicrobial Chemotherapy, 2017, 72, 1784-1793.	3.0	57
26	The risk and clinical outcome of candidemia depending on underlying malignancy. Intensive Care Medicine, 2017, 43, 652-662.	8.2	92
27	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
28	Diversity of Pneumocystis jirovecii Across Europe: A Multicentre Observational Study. EBioMedicine, 2017, 22, 155-163.	6.1	20
29	Fluconazole and Echinocandin Resistance of Candida glabrata Correlates Better with Antifungal Drug Exposure Rather than with MSH2 Mutator Genotype in a French Cohort of Patients Harboring Low Rates of Resistance. Frontiers in Microbiology, 2016, 7, 2038.	3.5	59
30	Recurrent episodes of Candidemia due to Candida glabrata, Candida tropicalis and Candida albicans with acquired echinocandin resistance. Medical Mycology Case Reports, 2016, 14, 20-23.	1.3	17
31	Novel Taxa Associated with Human Fungal Black-Grain Mycetomas: Emarellia grisea gen. nov., sp. nov., and Emarellia paragrisea sp. nov. Journal of Clinical Microbiology, 2016, 54, 1738-1745.	3.9	33
32	Environmental distribution of <i>Cryptococcus neoformans </i> and <i>C. gattii </i> around the Mediterranean basin. FEMS Yeast Research, 2016, 16, fow 045.	2.3	57
33	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
34	Typing Candida Species Using Microsatellite Length Polymorphism and Multilocus Sequence Typing. Methods in Molecular Biology, 2016, 1356, 199-214.	0.9	29
35	International Society of Human and Animal Mycology (ISHAM)-ITS reference DNA barcoding databaseâ€"the quality controlled standard tool for routine identification of human and animal pathogenic fungi. Medical Mycology, 2015, 53, 313-337.	0.7	252
36	Cryptococcosis Serotypes Impact Outcome and Provide Evidence of Cryptococcus neoformans Speciation. MBio, 2015, 6, e00311.	4.1	67

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37	Application of Isothermal Amplification Techniques for Identification of Madurella mycetomatis, the Prevalent Agent of Human Mycetoma. Journal of Clinical Microbiology, 2015, 53, 3280-3285.	3.9	36
38	Multicenter Outbreak of Infections by $\mbox{\sc i} \times \mbox{\sc Saprochaete clavata} <\mbox{\sc i} \times \mbox{\sc saprochaete clavata} <\mbox{\sc i} \times \mbox{\sc saprochaete clavata} <\mbox{\sc liv} \times \mbox{\sc liv} \times \mbox{\sc saprochaete clavata} <\mbox{\sc liv} \times \mbox{\sc liv} \times \times \mbox{\sc liv} \times \$	4.1	75
39	Misidentification of Saprochaete clavata as Magnusiomyces capitatus in Clinical Isolates: Utility of Internal Transcribed Spacer Sequencing and Matrix-Assisted Laser Desorption Ionization–Time of Flight Mass Spectrometry and Importance of Reliable Databases. Journal of Clinical Microbiology, 2014, 52, 2196-2198.	3.9	37
40	<i>Candida</i> spp. with Acquired Echinocandin Resistance, France, 2004–20101. Emerging Infectious Diseases, 2012, 18, 86-90.	4.3	116
41	Genetic Diversity and Genomic Plasticity of <i>Cryptococcus neoformans</i> AD Hybrid Strains. G3: Genes, Genomes, Genetics, 2012, 2, 83-97.	1.8	73
42	Antifungal Susceptibility Profiles of 1698 Yeast Reference Strains Revealing Potential Emerging Human Pathogens. PLoS ONE, 2012, 7, e32278.	2.5	25
43	Dynamics of Cryptococcus neoformans-Macrophage Interactions Reveal that Fungal Background Influences Outcome during Cryptococcal Meningoencephalitis in Humans. MBio, 2011, 2, .	4.1	102
44	Recent Exposure to Caspofungin or Fluconazole Influences the Epidemiology of Candidemia: a Prospective Multicenter Study Involving 2,441 Patients. Antimicrobial Agents and Chemotherapy, 2011, 55, 532-538.	3.2	294
45	Development of Echinocandin Resistance in Clavispora lusitaniae during Caspofungin Treatment. Journal of Clinical Microbiology, 2011, 49, 2304-2306.	3.9	34
46	Prior Caspofungin Exposure in Patients with Hematological Malignancies Is a Risk Factor for Subsequent Fungemia Due to Decreased Susceptibility in Candida spp.: a Case-Control Study in Paris, France. Antimicrobial Agents and Chemotherapy, 2011, 55, 5358-5361.	3.2	39
47	Mixed Infections and $\langle i \rangle$ In Vivo $\langle i \rangle$ Evolution in the Human Fungal Pathogen Cryptococcus neoformans. MBio, 2010, 1, .	4.1	88
48	Evidence That Graftâ€6ite Candidiasis after Kidney Transplantation Is Acquired during Organ Recovery: A Multicenter Study in France. Clinical Infectious Diseases, 2009, 48, 194-202.	5 <b>.</b> 8	105
49	<i>Debaryomyces hansenii</i> ( <i>Candida famata</i> ), a Rare Human Fungal Pathogen Often Misidentified as <i>Pichia guilliermondii</i> ( <i>Candida guilliermondii</i> ). Journal of Clinical Microbiology, 2008, 46, 3237-3242.	3.9	128
50	Detection of Caspofungin Resistance in <i>Candida</i> spp. by Etest. Journal of Clinical Microbiology, 2008, 46, 2389-2392.	3.9	31
51	Mutations in the <i>fks1</i> Gene in <i>Candida albicans</i> , <i>C. tropicalis</i> , and <i>C. krusei</i> Correlate with Elevated Caspofungin MICs Uncovered in AM3 Medium Using the Method of the European Committee on Antibiotic Susceptibility Testing. Antimicrobial Agents and Chemotherapy, 2008, 52, 3092-3098.	3.2	123
52	Clonal Population of Flucytosine-Resistant <i>Candida tropicalis</i> France. Emerging Infectious Diseases, 2008, 14, 557-565.	4.3	50
53	Comparison of Microsatellite Length Polymorphism and Multilocus Sequence Typing for DNA-Based Typing of <i>Candida albicans</i> . Journal of Clinical Microbiology, 2007, 45, 3958-3963.	3.9	51
54	Acquired resistance to echinocandins in Candida albicans: case report and review. Journal of Antimicrobial Chemotherapy, 2007, 59, 1076-1083.	3.0	136

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55	Molecular Identification of Black-Grain Mycetoma Agents. Journal of Clinical Microbiology, 2006, 44, 3517-3523.	3.9	89