MarÃ-a Guadalupe FrÃ-as De LeÃ³n

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Detection and Molecular Identification of Eight Candida Species in Clinical Samples by Simplex PCR. Microorganisms, 2022, 10, 374.	3.6	6
2	Selection of Polymorphic Patterns Obtained by RAPD-PCR through Qualitative and Quantitative Analyses to Differentiate Aspergillus fumigatus. Journal of Fungi (Basel, Switzerland), 2022, 8, 296.	3.5	2
3	Vaccines Against COVID-19: A Review. Vaccines, 2022, 10, 414.	4.4	8

Antifungal Resistance in Clinical Isolates of Candida glabrata in Ibero-America. Journal of Fungi (Basel,) Tj ETQq0 0 0.rgBT /Overlock 10 Tr

5	Diagnostic accuracy of antigen detection in urine and molecular assays testing in different clinical samples for the diagnosis of progressive disseminated histoplasmosis in patients living with HIV/AIDS: A prospective multicenter study in Mexico. PLoS Neglected Tropical Diseases, 2021, 15, e0009215.	3.0	25
6	Rhinocerebral mucormycosis to the rise? The impact of the worldwide diabetes epidemic. Anais Brasileiros De Dermatologia, 2021, 96, 196-199.	1.1	8
7	Molecular identification of yeasts from the order Trichosporonales causing superficial infections. Revista Iberoamericana De Micologia, 2021, 38, 119-124.	0.9	6
8	Epidemiology of Systemic Mycoses in the COVID-19 Pandemic. Journal of Fungi (Basel, Switzerland), 2021, 7, 556.	3.5	23
9	Candida glabrata Antifungal Resistance and Virulence Factors, a Perfect Pathogenic Combination. Pharmaceutics, 2021, 13, 1529.	4.5	17
10	Phylogenetic Identification, Diversity, and Richness of Aspergillus from Homes in Havana, Cuba. Microorganisms, 2021, 9, 115.	3.6	6
11	Antifungal Resistance in Candida auris: Molecular Determinants. Antibiotics, 2020, 9, 568.	3.7	38
12	Identification of CSP Types and Genotypic Variability of Clinical and Environmental Isolates of Aspergillus fumigatus from Different Geographic Origins. Microorganisms, 2020, 8, 688.	3.6	7
13	Usefulness of a multiplex PCR for the rapid identification of Candida glabrata species complex in Mexican clinical isolates. Revista Do Instituto De Medicina Tropical De Sao Paulo, 2019, 61, e37.	1.1	3
14	Chromoblastomycosis caused by Rhinocladiella aquaspersa: first case report in Guatemala. Anais Brasileiros De Dermatologia, 2019, 94, 574-577.	1.1	7
15	Genotyping of Leptospira interrogans isolates from Mexican patients. Revista Do Instituto De Medicina Tropical De Sao Paulo, 2019, 61, e26.	1.1	0
16	Update of Vulvovaginal Candidiasis in Pregnant and Non-pregnant Patients. Current Fungal Infection Reports, 2019, 13, 181-190.	2.6	1
17	Histoplasmosis: diagnostic challenges. Case Reports, 2019, 5, 85-88.	0.0	0
18	Identification of AspergillusÂtubingensis in a primary skin infection. Journal De Mycologie Medicale, 2018, 28, 274-278.	1.5	16

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#	Article	IF	CITATIONS
19	Selection of Specific Peptides for Coccidioides spp. Obtained from Antigenic Fractions through SDS-PAGE and Western Blot Methods by the Recognition of Sera from Patients with Coccidioidomycosis. Molecules, 2018, 23, 3145.	3.8	6
20	Onychomycosis Due to Aspergillus spp.: a Current Review. Current Fungal Infection Reports, 2018, 12, 112-119.	2.6	1
21	Current status of the etiology of candidiasis in Mexico. Revista Iberoamericana De Micologia, 2017, 34, 203-210.	0.9	19
22	Usefulness of molecular markers in the diagnosis of occupational and recreational histoplasmosis outbreaks. Folia Microbiologica, 2017, 62, 111-116.	2.3	12
23	The habitat of Coccidioides spp. and the role of animals as reservoirs and disseminators in nature. BMC Infectious Diseases, 2016, 16, 550.	2.9	46
24	AFLP analysis reveals high genetic diversity but low population structure in Coccidioides posadasiiisolates from Mexico and Argentina. BMC Infectious Diseases, 2013, 13, 411.	2.9	22
25	Development of Specific Sequence-Characterized Amplified Region Markers for Detecting Histoplasma capsulatum in Clinical and Environmental Samples. Journal of Clinical Microbiology, 2012, 50, 673-679.	3.9	24
26	Phenotypic characteristics of isolates of Aspergillus section Fumigatifrom different geographic origins and their relationships with genotypic characteristics. BMC Infectious Diseases, 2011, 11, 116.	2.9	11