

Lalitha Gade

List of Publications by Year in descending order

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Version: 2024-02-01

29
papers

2,236
citations

361413

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477307

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docs citations

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times ranked

1840
citing authors

#	ARTICLE	IF	CITATIONS
1	Genomic insights into multidrug-resistance, mating and virulence in <i>Candida auris</i> and related emerging species. <i>Nature Communications</i> , 2018, 9, 5346.	12.8	298
2	Necrotizing Cutaneous Mucormycosis after a Tornado in Joplin, Missouri, in 2011. <i>New England Journal of Medicine</i> , 2012, 367, 2214-2225.	27.0	297
3	Tracing the Evolutionary History and Global Expansion of <i>Candida auris</i> Using Population Genomic Analyses. <i>MBio</i> , 2020, 11, .	4.1	224
4	Multiple introductions and subsequent transmission of multidrug-resistant <i>Candida auris</i> in the USA: a molecular epidemiological survey. <i>Lancet Infectious Diseases</i> , The, 2018, 18, 1377-1384.	9.1	204
5	Valley Fever: Finding New Places for an Old Disease: <i>Coccidioides immitis</i> Found in Washington State Soil Associated With Recent Human Infection. <i>Clinical Infectious Diseases</i> , 2015, 60, e1-e3.	5.8	153
6	Molecular Epidemiology of <i>Candida auris</i> in Colombia Reveals a Highly Related, Countrywide Colonization With Regional Patterns in Amphotericin B Resistance. <i>Clinical Infectious Diseases</i> , 2019, 68, 15-21.	5.8	132
7	Notes from the Field: Ongoing Transmission of <i>Candida auris</i> in Health Care Facilities – United States, June 2016–May 2017. <i>Morbidity and Mortality Weekly Report</i> , 2017, 66, 514-515.	15.1	124
8	Mutations in <i>TAC1B</i> : a Novel Genetic Determinant of Clinical Fluconazole Resistance in <i>Candida auris</i> . <i>MBio</i> , 2020, 11, .	4.1	101
9	Utility of (1 ³ C)-β-D-Glucan Testing for Diagnostics and Monitoring Response to Treatment During the Multistate Outbreak of Fungal Meningitis and Other Infections. <i>Clinical Infectious Diseases</i> , 2014, 58, 622-630.	5.8	85
10	Local Population Structure and Patterns of Western Hemisphere Dispersal for <i>Coccidioides</i> spp., the Fungal Cause of Valley Fever. <i>MBio</i> , 2016, 7, e00550-16.	4.1	71
11	Whole Genome Sequence Typing to Investigate the <i>Apophysomyces</i> Outbreak following a Tornado in Joplin, Missouri, 2011. <i>PLoS ONE</i> , 2012, 7, e49989.	2.5	66
12	Detection of Fungal DNA in Human Body Fluids and Tissues during a Multistate Outbreak of Fungal Meningitis and Other Infections. <i>Eukaryotic Cell</i> , 2013, 12, 677-683.	3.4	62
13	Clade-specific chromosomal rearrangements and loss of subtelomeric adhesins in <i>Candida auris</i> . <i>Genetics</i> , 2021, 218, .	2.9	54
14	Whole-Genome Analysis of <i>Exserohilum rostratum</i> from an Outbreak of Fungal Meningitis and Other Infections. <i>Journal of Clinical Microbiology</i> , 2014, 52, 3216-3222.	3.9	52
15	Preliminary Laboratory Report of Fungal Infections Associated with Contaminated Methylprednisolone Injections. <i>Journal of Clinical Microbiology</i> , 2013, 51, 2654-2661.	3.9	41
16	<i>Coccidioides immitis</i> identified in soil outside of its known range - Washington, 2013. <i>Morbidity and Mortality Weekly Report</i> , 2014, 63, 450.	15.1	30
17	Utility of Whole-Genome Sequencing to Ascertain Locally Acquired Cases of <i>Coccidioidomycosis</i> , Washington, USA. <i>Emerging Infectious Diseases</i> , 2019, 25, 501-506.	4.3	24
18	Understanding the Emergence of Multidrug-Resistant <i>Candida</i> : Using Whole-Genome Sequencing to Describe the Population Structure of <i>Candida haemulonii</i> Species Complex. <i>Frontiers in Genetics</i> , 2020, 11, 554.	2.3	24

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19	Detection of mucormycetes and other pathogenic fungi in formalin fixed paraffin embedded and fresh tissues using the extended region of 28S rDNA. <i>Medical Mycology</i> , 2016, 55, myw083.	0.7	23
20	The detection of <i>Coccidioides</i> from ambient air in Phoenix, Arizona: Evidence of uneven distribution and seasonality. <i>Medical Mycology</i> , 2020, 58, 552-559.	0.7	23
21	<i>Candida auris</i> outbreak involving liver transplant recipients in a surgical intensive care unit. <i>American Journal of Transplantation</i> , 2020, 20, 3673-3679.	4.7	23
22	Utility of Real-Time PCR for Detection of <i>Exserohilum rostratum</i> in Body and Tissue Fluids during the Multistate Outbreak of Fungal Meningitis and Other Infections. <i>Journal of Clinical Microbiology</i> , 2015, 53, 618-625.	3.9	20
23	Genomic Diversity of Azole-Resistant <i>Aspergillus fumigatus</i> in the United States. <i>MBio</i> , 2021, 12, e0180321.	4.1	17
24	Rapid Assessment and Containment of <i>Candida auris</i> Transmission in Postacute Care Settings—Orange County, California, 2019. <i>Annals of Internal Medicine</i> , 2021, 174, 1554-1562.	3.9	17
25	Factors Influencing Distribution of <i>Coccidioides immitis</i> in Soil, Washington State, 2016. <i>MSphere</i> , 2021, 6, e0059821.	2.9	17
26	MycoSNP: A Portable Workflow for Performing Whole-Genome Sequencing Analysis of <i>Candida auris</i> . <i>Methods in Molecular Biology</i> , 2022, , 215-228.	0.9	13
27	Genome Sequence of a Multidrug-Resistant <i>Candida haemulonii</i> Isolate from a Patient with Chronic Leg Ulcers in Israel. <i>Genome Announcements</i> , 2018, 6, .	0.8	11
28	<i>Rhizopus microsporus</i> Infections Associated with Surgical Procedures, Argentina, 2006–2014. <i>Emerging Infectious Diseases</i> , 2020, 26, 937-944.	4.3	11
29	Genome Sequence of the Amphotericin B-Resistant <i>Candida duobushaemulonii</i> Strain B09383. <i>Genome Announcements</i> , 2018, 6, .	0.8	9