

Julio Guerra

List of Publications by Year in descending order

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Version: 2024-02-01

18
papers

176
citations

1307594

7
h-index

1125743

13
g-index

20
all docs

20
docs citations

20
times ranked

285
citing authors

#	ARTICLE	IF	CITATIONS
1	Heterogeneous fitness landscape cues, <i>pknG</i> high expression, and phthiocerol dimycocerosate low production of <i>Mycobacterium tuberculosis</i> ATCC25618 <i>rpoB</i> S450L in enriched broth. <i>Tuberculosis</i> , 2022, 132, 102156.	1.9	3
2	Close Related Drug-Resistance Beijing Isolates of <i>Mycobacterium tuberculosis</i> Reveal a Different Transcriptomic Signature in a Murine Disease Progression Model. <i>International Journal of Molecular Sciences</i> , 2022, 23, 5157.	4.1	3
3	First approach to the population structure of <i>Mycobacterium tuberculosis</i> complex in the indigenous population in Puerto Nariño-Amazonas, Colombia. <i>PLoS ONE</i> , 2021, 16, e0245084.	2.5	5
4	Profiling the immune response to <i>Mycobacterium tuberculosis</i> Beijing family infection: a perspective from the transcriptome. <i>Virulence</i> , 2021, 12, 1689-1704.	4.4	9
5	Title is missing!. , 2021, 16, e0245084.		0
6	Title is missing!. , 2021, 16, e0245084.		0
7	Title is missing!. , 2021, 16, e0245084.		0
8	Title is missing!. , 2021, 16, e0245084.		0
9	Genetic diversity of <i>Mycobacterium tuberculosis</i> clinical isolates from HIV-TB patients from two public hospitals at Bogotá, Colombia. <i>Infection, Genetics and Evolution</i> , 2020, 77, 104059.	2.3	0
10	Population structure of multidrug-resistant <i>Mycobacterium tuberculosis</i> clinical isolates in Colombia. <i>Tuberculosis</i> , 2020, 125, 102011.	1.9	4
11	Characterization of clinical isolates of <i>Mycobacterium tuberculosis</i> from indigenous peoples of Colombia. <i>Biomedica</i> , 2019, 39, 78-92.	0.7	3
12	Active and latent tuberculosis among inmates in La Esperanza prison in Guaduas, Colombia. <i>PLoS ONE</i> , 2019, 14, e0209895.	2.5	12
13	Circulation of <i>M. tuberculosis</i> Beijing genotype in Latin America and the Caribbean. <i>Pathogens and Global Health</i> , 2019, 113, 336-351.	2.3	13
14	Comparative genomic analysis of <i>Mycobacterium tuberculosis</i> Beijing-like strains revealed specific genetic variations associated with virulence and drug resistance. <i>Infection, Genetics and Evolution</i> , 2017, 54, 314-323.	2.3	30
15	Complete Genome Sequence of the Clinical Beijing-Like Strain <i>Mycobacterium tuberculosis</i> 323 Using the PacBio Real-Time Sequencing Platform. <i>Genome Announcements</i> , 2015, 3, .	0.8	10
16	Virulence and Immune Response Induced by <i>Mycobacterium avium</i> Complex Strains in a Model of Progressive Pulmonary Tuberculosis and Subcutaneous Infection in BALB/c Mice. <i>Infection and Immunity</i> , 2013, 81, 4001-4012.	2.2	18
17	A first insight on the population structure of <i>Mycobacterium tuberculosis</i> complex as studied by spoligotyping and MIRU-VNTRs in Bogotá, Colombia. <i>Infection, Genetics and Evolution</i> , 2012, 12, 657-663.	2.3	42
18	First case of multidrug-resistant tuberculosis caused by a rare "Beijing-like" genotype of <i>Mycobacterium tuberculosis</i> in Bogotá, Colombia. <i>Infection, Genetics and Evolution</i> , 2010, 10, 678-681.	2.3	16