

# Sidra E G Ezidio Gonçalves Vasconcello

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/949030/publications.pdf>

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10  
papers

101  
citations

1307594

7  
h-index

1372567

10  
g-index

10  
all docs

10  
docs citations

10  
times ranked

212  
citing authors

#	ARTICLE	IF	CITATIONS
1	Molecular Typing of Mycobacterium bovis from Cattle Reared in Midwest Brazil. PLoS ONE, 2016, 11, e0162459.	2.5	24
2	Genetic diversity of Mycobacterium tuberculosis isoniazid mono-resistant and multidrug-resistant in Rio Grande do Sul, a tuberculosis high-burden state in Brazil. Tuberculosis, 2018, 110, 36-43.	1.9	16
3	Sequence and structural characterization of tbnat gene in isoniazid-resistant Mycobacterium tuberculosis: Identification of new mutations. Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis, 2011, 712, 33-39.	1.0	10
4	Genetic Clustering of Tuberculosis in an Indigenous Community of Brazil. American Journal of Tropical Medicine and Hygiene, 2018, 98, 372-375.	1.4	10
5	Genetic diversity of Mycobacterium tuberculosis from Pará, Brazil, reveals a higher frequency of ancestral strains than previously reported in South America. Infection, Genetics and Evolution, 2017, 56, 62-72.	2.3	9
6	Insights into the population structure of Mycobacterium tuberculosis using spoligotyping and RDRio in a southeastern Brazilian prison unit. Infection, Genetics and Evolution, 2014, 26, 194-202.	2.3	8
7	Frequency of the Mycobacterium tuberculosis RDRio genotype and its association with multidrug-resistant tuberculosis. BMC Infectious Diseases, 2019, 19, 556.	2.9	8
8	Antiretroviral therapy-induced paradoxical worsening of previously healed Mycobacterium haemophilum cutaneous lesions in advanced HIV infection. Revista Do Instituto De Medicina Tropical De Sao Paulo, 2019, 61, e71.	1.1	8
9	Whole genome sequence of Mycobacterium kansasii isolates of the genotype 1 from Brazilian patients with pulmonary disease demonstrates considerable heterogeneity. Memórias Do Instituto Oswaldo Cruz, 2018, 113, e180085.	1.6	7
10	Genome Comparison of an Ancestral Isolate and a Modern Isolate of Mycobacterium tuberculosis of the Beijing Lineage from São Paulo, Brazil. Genome Announcements, 2015, 3, .	0.8	1