

# Serej D Ley

## List of Publications by Year in descending order

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Version: 2024-02-01

12  
papers

529  
citations

1163117

8  
h-index

1281871

11  
g-index

13  
all docs

13  
docs citations

13  
times ranked

990  
citing authors

#	ARTICLE	IF	CITATIONS
1	Transmission, distribution and drug resistance-conferring mutations of extensively drug-resistant tuberculosis in the Western Cape Province, South Africa. <i>Microbial Genomics</i> , 2022, 8, .	2.0	3
2	Local adaptation in populations of <i>Mycobacterium tuberculosis</i> endemic to the Indian Ocean Rim. <i>F1000Research</i> , 2021, 10, 60.	1.6	13
3	Melting the <i>eis</i> : Nondetection of Kanamycin Resistance Markers by Routine Diagnostic Tests and Identification of New <i>eis</i> Promoter Variants. <i>Antimicrobial Agents and Chemotherapy</i> , 2021, 65, e0250220.	3.2	0
4	Multiple Introductions of <i>Mycobacterium tuberculosis</i> Lineage 2“Beijing Into Africa Over Centuries. <i>Frontiers in Ecology and Evolution</i> , 2019, 7, .	2.2	29
5	Detection of Second Line Drug Resistance among Drug Resistant <i>Mycobacterium Tuberculosis</i> Isolates in Botswana. <i>Pathogens</i> , 2019, 8, 208.	2.8	9
6	Bedaquiline Microheteroresistance after Cessation of Tuberculosis Treatment. <i>New England Journal of Medicine</i> , 2019, 380, 2178-2180.	27.0	52
7	Genetic diversity of <i>Mycobacterium tuberculosis</i> strains circulating in Botswana. <i>PLoS ONE</i> , 2019, 14, e0216306.	2.5	9
8	Deciphering Within-Host Microevolution of <i>Mycobacterium tuberculosis</i> through Whole-Genome Sequencing: the Phenotypic Impact and Way Forward. <i>Microbiology and Molecular Biology Reviews</i> , 2019, 83, .	6.6	43
9	<i>Mycobacterium tuberculosis</i> lineage 4 comprises globally distributed and geographically restricted sublineages. <i>Nature Genetics</i> , 2016, 48, 1535-1543.	21.4	326
10	Diversity of <i>Mycobacterium tuberculosis</i> and drug resistance in different provinces of Papua New Guinea. <i>BMC Microbiology</i> , 2014, 14, 307.	3.3	19
11	Tuberculosis in Papua New Guinea: from yesterday until today. <i>Microbes and Infection</i> , 2014, 16, 607-614.	1.9	17
12	A microarray-based system for the simultaneous analysis of single nucleotide polymorphisms in human genes involved in the metabolism of anti-malarial drugs. <i>Malaria Journal</i> , 2009, 8, 285.	2.3	7