

Timothy C Rodwell

List of Publications by Year in descending order

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Version: 2024-02-01

35
papers

1,915
citations

430874

18
h-index

361022

35
g-index

40
all docs

40
docs citations

40
times ranked

2278
citing authors

#	ARTICLE	IF	CITATIONS
1	Prediction of Susceptibility to First-Line Tuberculosis Drugs by DNA Sequencing. <i>New England Journal of Medicine</i> , 2018, 379, 1403-1415.	27.0	405
2	A standardised method for interpreting the association between mutations and phenotypic drug resistance in <i>Mycobacterium tuberculosis</i> . <i>European Respiratory Journal</i> , 2017, 50, 1701354.	6.7	273
3	Genetic Mutations Associated with Isoniazid Resistance in <i>Mycobacterium tuberculosis</i> : A Systematic Review. <i>PLoS ONE</i> , 2015, 10, e0119628.	2.5	236
4	The 2021 WHO catalogue of <i>Mycobacterium tuberculosis</i> complex mutations associated with drug resistance: a genotypic analysis. <i>Lancet Microbe</i> , The, 2022, 3, e265-e273.	7.3	114
5	Predicting Extensively Drug-Resistant <i>Mycobacterium tuberculosis</i> Phenotypes with Genetic Mutations. <i>Journal of Clinical Microbiology</i> , 2014, 52, 781-789.	3.9	99
6	Novel <i>katG</i> mutations causing isoniazid resistance in clinical <i>M. tuberculosis</i> isolates. <i>Emerging Microbes and Infections</i> , 2015, 4, 1-9.	6.5	95
7	Detection of Low-Level Mixed-Population Drug Resistance in <i>Mycobacterium tuberculosis</i> Using High Fidelity Amplicon Sequencing. <i>PLoS ONE</i> , 2015, 10, e0126626.	2.5	93
8	Rapid Drug Susceptibility Testing of Drug-Resistant <i>Mycobacterium tuberculosis</i> Isolates Directly from Clinical Samples by Use of Amplicon Sequencing: a Proof-of-Concept Study. <i>Journal of Clinical Microbiology</i> , 2016, 54, 2058-2067.	3.9	76
9	Integrating standardized whole genome sequence analysis with a global <i>Mycobacterium tuberculosis</i> antibiotic resistance knowledgebase. <i>Scientific Reports</i> , 2018, 8, 15382.	3.3	75
10	Application of Targeted Next-Generation Sequencing Assay on a Portable Sequencing Platform for Culture-Free Detection of Drug-Resistant Tuberculosis from Clinical Samples. <i>Journal of Clinical Microbiology</i> , 2020, 58, .	3.9	57
11	Whole-genome and targeted sequencing of drug-resistant <i>Mycobacterium tuberculosis</i> on the iSeq100 and MiSeq: A performance, ease-of-use, and cost evaluation. <i>PLoS Medicine</i> , 2019, 16, e1002794.	8.4	49
12	Correlating <i>rrs</i> and <i>eis</i> promoter mutations in clinical isolates of <i>Mycobacterium tuberculosis</i> with phenotypic susceptibility levels to the second-line injectables. <i>International Journal of Mycobacteriology</i> , 2016, 5, 1-6.	0.6	42
13	Evaluation of Pyrosequencing for Detecting Extensively Drug-Resistant <i>Mycobacterium tuberculosis</i> among Clinical Isolates from Four High-Burden Countries. <i>Antimicrobial Agents and Chemotherapy</i> , 2015, 59, 414-420.	3.2	36
14	Correlating Minimum Inhibitory Concentrations of ofloxacin and moxifloxacin with <i>gyrA</i> mutations using the genotype MTBDRsl assay. <i>Tuberculosis</i> , 2015, 95, 137-141.	1.9	34
15	Performance of the Xpert MTB/RIF assay for the diagnosis of pulmonary tuberculosis and rifampin resistance in a low-incidence, high-resource setting. <i>PLoS ONE</i> , 2017, 12, e0186139.	2.5	33
16	Requiring smartphone ownership for mHealth interventions: who could be left out?. <i>BMC Public Health</i> , 2020, 20, 81.	2.9	31
17	The Global Consortium for Drug-resistant Tuberculosis Diagnostics (GCDD): design of a multi-site, head-to-head study of three rapid tests to detect extensively drug-resistant tuberculosis. <i>Trials</i> , 2014, 15, 434.	1.6	28
18	Defining multidrug-resistant tuberculosis: correlating GenoType MTBDR plus assay results with minimum inhibitory concentrations. <i>Diagnostic Microbiology and Infectious Disease</i> , 2015, 82, 49-53.	1.8	21

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19	Increased Tuberculosis Patient Mortality Associated with Mycobacterium tuberculosis Mutations Conferring Resistance to Second-Line Antituberculous Drugs. <i>Journal of Clinical Microbiology</i> , 2017, 55, 1928-1937.	3.9	16
20	Next-generation sequencing-based user-friendly platforms for drug-resistant tuberculosis diagnosis: A promise for the near future. <i>International Journal of Mycobacteriology</i> , 2016, 5, S27-S28.	0.6	14
21	Distinct blood transcriptomic signature of treatment in latent tuberculosis infected individuals at risk of developing active disease. <i>Tuberculosis</i> , 2021, 131, 102127.	1.9	13
22	Impact of Fluoroquinolone Use on Mortality Among a Cohort of Patients With Suspected Drug-Resistant Tuberculosis. <i>Clinical Infectious Diseases</i> , 2017, 65, 772-778.	5.8	12
23	Shedding light on the performance of a pyrosequencing assay for drug-resistant tuberculosis diagnosis. <i>BMC Infectious Diseases</i> , 2016, 16, 458.	2.9	9
24	Detection and quantification of Mycobacterium tuberculosis antigen CFP10 in serum and urine for the rapid diagnosis of active tuberculosis disease. <i>Scientific Reports</i> , 2021, 11, 19193.	3.3	8
25	Surveillance or support: The experience of direct observation during tuberculosis treatment. <i>Global Public Health</i> , 2018, 13, 804-818.	2.0	7
26	Cost analysis of rapid diagnostics for drug-resistant tuberculosis. <i>BMC Infectious Diseases</i> , 2018, 18, 102.	2.9	6
27	A novel technique for aging male African elephants (<i>Loxodonta africana</i>) using craniofacial photogrammetry and geometric morphometrics. <i>Mammalian Biology</i> , 0, , .	1.5	5
28	Variants in Bedaquiline-Candidate-Resistance Genes: Prevalence in Bedaquiline-Naive Patients, Effect on MIC, and Association with Mycobacterium tuberculosis Lineage. <i>Antimicrobial Agents and Chemotherapy</i> , 2022, 66, .	3.2	5
29	Using Mycobacterium tuberculosis Single-Nucleotide Polymorphisms To Predict Fluoroquinolone Treatment Response. <i>Antimicrobial Agents and Chemotherapy</i> , 2019, 63, .	3.2	4
30	Laboratory Evaluation of a Lateral-Flow Cell for Molecular Detection of First-Line and Second-Line Antituberculosis Drug Resistance. <i>Journal of Clinical Microbiology</i> , 2020, 58, .	3.9	3
31	Review of automated DNA extraction systems for sequencing-based solutions for drug-resistant tuberculosis detection. <i>Diagnostic Microbiology and Infectious Disease</i> , 2020, 98, 115096.	1.8	3
32	Assessing COVID-19-Related Knowledge, Attitudes, and Practices Among Hispanic Primary Care Patients: Protocol for a Cross-sectional Survey Study. <i>JMIR Research Protocols</i> , 2021, 10, e25265.	1.0	3
33	Rapid Detection of Extensively Drug-Resistant Tuberculosis in Clinical Samples Using a Novel Tabletop Platform: Protocol for a Prospective Clinical Study. <i>JMIR Research Protocols</i> , 2021, 10, e26748.	1.0	2
34	Knowledge, Attitude, Practices, and Vaccine Hesitancy Among the Latinx Community in Southern California Early in the COVID-19 Pandemic: Cross-sectional Survey. <i>JMIR Formative Research</i> , 2022, 6, e38351.	1.4	2
35	Detecting rifampin and isoniazid resistance in Mycobacterium tuberculosis direct from patient sputum using an automated integrated system. <i>Journal of Clinical Tuberculosis and Other Mycobacterial Diseases</i> , 2022, 27, 100304.	1.3	1