

Ruwen Jou

List of Publications by Year in descending order

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Version: 2024-02-01

29
papers

586
citations

759233

12
h-index

610901

24
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30
all docs

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docs citations

30
times ranked

900
citing authors

#	ARTICLE	IF	CITATIONS
1	Bedaquiline Drug Resistance Emergence Assessment in Multidrug-Resistant Tuberculosis (MDR-TB): a 5-Year Prospective <i>In Vitro</i> Surveillance Study of Bedaquiline and Other Second-Line Drug Susceptibility Testing in MDR-TB Isolates. <i>Journal of Clinical Microbiology</i> , 2022, 60, JCM0291920.	3.9	20
2	Bovine tuberculosis in Taiwan, 2008–2019. <i>Transboundary and Emerging Diseases</i> , 2022, 69, .	3.0	2
3	Molecular Epidemiological Characteristics of Mycobacterium abscessus Complex Derived from Non-Cystic Fibrosis Patients in Japan and Taiwan. <i>Microbiology Spectrum</i> , 2022, 10, e0057122.	3.0	7
4	Disputed <i>rpoB</i> Mutations in Mycobacterium tuberculosis and Tuberculosis Treatment Outcomes. <i>Antimicrobial Agents and Chemotherapy</i> , 2021, 65, e0157320.	3.2	7
5	Sensible Functional Linear Discriminant Analysis Effectively Discriminates Enhanced Raman Spectra of <i>Mycobacterium</i> Species. <i>Analytical Chemistry</i> , 2021, 93, 2785-2792.	6.5	15
6	Primary Bedaquiline Resistance Among Cases of Drug-Resistant Tuberculosis in Taiwan. <i>Frontiers in Microbiology</i> , 2021, 12, 754249.	3.5	19
7	A new oligonucleotide array for the detection of multidrug and extensively drug-resistance tuberculosis. <i>Scientific Reports</i> , 2019, 9, 4425.	3.3	10
8	Impact of universal drug susceptibility testing and effective management of multidrug-resistant tuberculosis in Taiwan. <i>PLoS ONE</i> , 2019, 14, e0214792.	2.5	4
9	Redefining MDR-TB: Comparison of Mycobacterium tuberculosis clinical isolates from Russia and Taiwan. <i>Infection, Genetics and Evolution</i> , 2019, 72, 141-146.	2.3	10
10	Performance of an Xpert-based diagnostic algorithm for the rapid detection of drug-resistant tuberculosis among high-risk populations in a low-incidence setting. <i>PLoS ONE</i> , 2018, 13, e0200755.	2.5	14
11	Susceptibilities of MDR Mycobacterium tuberculosis isolates to unconventional drugs compared with their reported pharmacokinetic/pharmacodynamic parameters. <i>Journal of Antimicrobial Chemotherapy</i> , 2017, 72, 1678-1687.	3.0	18
12	Using genotyping to delineate tuberculosis transmission in long-term care facilities: single facility 4-year experience. <i>BMC Infectious Diseases</i> , 2017, 17, 421.	2.9	2
13	CLEC9A modulates macrophage-mediated neutrophil recruitment in response to heat-killed Mycobacterium tuberculosis H37Ra. <i>PLoS ONE</i> , 2017, 12, e0186780.	2.5	11
14	Population-Based Drug Resistance Surveillance of Multidrug-Resistant Tuberculosis in Taiwan, 2007-2014. <i>PLoS ONE</i> , 2016, 11, e0165222.	2.5	12
15	An integrated MDR-TB management programme results in favourable outcomes in northern Taiwan. <i>European Respiratory Journal</i> , 2015, 45, 272-275.	6.7	7
16	Sputum culture conversion as a prognostic marker for end-of-treatment outcome in patients with multidrug-resistant tuberculosis: a secondary analysis of data from two observational cohort studies. <i>Lancet Respiratory Medicine</i> , 2015, 3, 201-209.	10.7	116
17	Leprosy in Taiwan, 2002–2011. <i>Journal of the Formosan Medical Association</i> , 2014, 113, 579-580.	1.7	5
18	Molecular Detection of Rifabutin-Susceptible Mycobacterium tuberculosis. <i>Journal of Clinical Microbiology</i> , 2012, 50, 2085-2088.	3.9	17

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19	Molecular diagnosis of childhood tuberculosis and infection with Bacilli Calmette-Guerin in Taiwan. Journal of the Formosan Medical Association, 2011, 110, 759-761.	1.7	2
20	Characteristics of multidrug-resistant Mycobacterium tuberculosis in Taiwan: A population-based study. Infection, Genetics and Evolution, 2011, 11, 633-639.	2.3	10
21	Evaluation of the Rapid MGIT TBc Identification Test for Culture Confirmation of Mycobacterium tuberculosis Complex Strain Detection. Journal of Clinical Microbiology, 2011, 49, 802-807.	3.9	51
22	Single nucleotide polymorphisms in cell wall biosynthesis-associated genes and phylogeny of Mycobacterium tuberculosis lineages. Infection, Genetics and Evolution, 2010, 10, 459-466.	2.3	10
23	Management of a Nosocomial Outbreak of <i>Mycobacterium tuberculosis</i> Beijing/W Genotype in Taiwan: an Emphasis on Case Tracing with High-Resolution Computed Tomography. Japanese Journal of Infectious Diseases, 2010, 63, 199-203.	1.2	4
24	Performance Assessment of the GenoType MTBDR <i>plus</i> Test and DNA Sequencing in Detection of Multidrug-Resistant <i>Mycobacterium tuberculosis</i> . Journal of Clinical Microbiology, 2009, 47, 2520-2524.	3.9	80
25	Spoligotypes of Mycobacterium tuberculosis isolates of a high tuberculosis burden aboriginal township in Taiwan. Infection, Genetics and Evolution, 2008, 8, 553-557.	2.3	10
26	Human Tuberculosis Caused by <i>Mycobacterium bovis</i> , Taiwan. Emerging Infectious Diseases, 2008, 14, 515-517.	4.3	16
27	Laboratory Investigation of a Nosocomial Transmission of Tuberculosis at a District General Hospital. Journal of the Formosan Medical Association, 2007, 106, 520-527.	1.7	7
28	Genetic Diversity of Multidrug-Resistant Mycobacterium tuberculosis Isolates and Identification of 11 Novel rpoB Alleles in Taiwan. Journal of Clinical Microbiology, 2005, 43, 1390-1394.	3.9	42
29	Distribution of the Beijing Family Genotypes of Mycobacterium tuberculosis in Taiwan. Journal of Clinical Microbiology, 2005, 43, 95-100.	3.9	58