

Jean-Luc Berland

List of Publications by Year in descending order

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Version: 2024-02-01

16
papers

375
citations

933447

10
h-index

996975

15
g-index

19
all docs

19
docs citations

19
times ranked

554
citing authors

#	ARTICLE	IF	CITATIONS
1	Rifampicin exposure reveals within-host <i>Mycobacterium tuberculosis</i> diversity in patients with delayed culture conversion. <i>PLoS Pathogens</i> , 2021, 17, e1009643.	4.7	10
2	Expanded tracking of a Beijing <i>Mycobacterium tuberculosis</i> strain involved in an outbreak in France. <i>Travel Medicine and Infectious Disease</i> , 2021, 44, 102167.	3.0	2
3	Routine survey of <i>Mycobacterium tuberculosis</i> isolates reveals nosocomial transmission. <i>European Respiratory Journal</i> , 2020, 55, 1901888.	6.7	6
4	Whole-genome sequencing in drug susceptibility testing of <i>Mycobacterium tuberculosis</i> in routine practice in Lyon, France. <i>International Journal of Antimicrobial Agents</i> , 2020, 55, 105912.	2.5	32
5	First insights into the genetic characteristics and drug resistance of <i>Mycobacterium tuberculosis</i> population collected during the first national tuberculosis prevalence survey of Lao PDR (2010–2011). <i>BMC Infectious Diseases</i> , 2019, 19, 851.	2.9	3
6	Subcultured <i>Mycobacterium tuberculosis</i> isolates on different growth media are fully representative of bacteria within clinical samples. <i>Tuberculosis</i> , 2019, 116, 61-66.	1.9	7
7	Prospective Whole-Genome Sequencing in Tuberculosis Outbreak Investigation, France, 2017–2018. <i>Emerging Infectious Diseases</i> , 2019, 25, 589-592.	4.3	25
8	High-Throughput <i>Mycobacterium tuberculosis</i> Interspersed Repetitive-Unit Variable-Number Tandem-Repeat Genotyping for <i>Mycobacterium tuberculosis</i> Epidemiological Studies. <i>Journal of Clinical Microbiology</i> , 2015, 53, 498-503.	3.9	17
9	Draft Genome Sequence of <i>Mycobacterium tuberculosis</i> Clinical Strain G-12-005. <i>Genome Announcements</i> , 2014, 2, .	0.8	0
10	Resistance of <i>Mycobacterium tuberculosis</i> to antibiotics in Lao PDR: first multicentric study conducted in 3 hospitals. <i>BMC Infectious Diseases</i> , 2013, 13, 275.	2.9	10
11	The Human Metapneumovirus Matrix Protein Stimulates the Inflammatory Immune Response In Vitro. <i>PLoS ONE</i> , 2011, 6, e17818.	2.5	13
12	Vaccine-induced early control of hepatitis C virus infection in chimpanzees fails to impact on hepatic PD-1 and chronicity. <i>Hepatology</i> , 2007, 45, 602-613.	7.3	84
13	False-Positive Results in a Recombinant Severe Acute Respiratory Syndrome-Associated Coronavirus (SARS-CoV) Nucleocapsid-Based Western Blot Assay Were Rectified by the Use of Two Subunits (S1 and N) Tj ETQq13110.7843134 rgBT /	1.0	0
14	Modulation of Vaccine-Induced Immune Responses to Hepatitis C Virus in Rhesus Macaques by Altering Priming before Adenovirus Boosting. <i>Journal of Infectious Diseases</i> , 2005, 192, 920-929.	4.0	38
15	Comparative immunogenicity analysis of modified vaccinia Ankara vectors expressing native or modified forms of hepatitis C virus E1 and E2 glycoproteins. <i>Vaccine</i> , 2004, 22, 3917-3928.	3.8	29
16	Antigenic relevance of F protein in chronic hepatitis C virus infection. <i>Hepatology</i> , 2004, 40, 900-909.	7.3	59