Valérie Bouchez

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5242440/publications.pdf

Version: 2024-02-01

840776 713466 21 661 11 21 citations h-index g-index papers 23 23 23 620 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Global spatial dynamics and vaccine-induced fitness changes of <i>Bordetella pertussis</i> Translational Medicine, 2022, 14, eabn3253.	12.4	22
2	A comprehensive resource for Bordetella genomic epidemiology and biodiversity studies. Nature Communications, 2022, 13, .	12.8	11
3	Ongoing diphtheria outbreak in Yemen: a cross-sectional and genomic epidemiology study. Lancet Microbe, The, 2021, 2, e386-e396.	7.3	26
4	Evolution of Bordetella pertussis over a 23-year period in France, 1996 to 2018. Eurosurveillance, 2021, 26, .	7.0	15
5	Corynebacterium rouxii sp. nov., a novel member of the diphtheriae species complex. Research in Microbiology, 2020, 171, 122-127.	2.1	45
6	<i>Bordetella parapertussis</i> Bacteremia: Clinical Expression and Bacterial Genomics. Open Forum Infectious Diseases, 2019, 6, ofz122.	0.9	10
7	Pertussis epidemiology in Tunisian infants and children and characterization of Bordetella pertussis isolates: results of a 9-year surveillance study, 2007 to 2016. Journal of Medical Microbiology, 2019, 68, 241-247.	1.8	14
8	Genome characteristics of Bordetella pertussis isolates from Tunisia. Journal of Medical Microbiology, 2019, 68, 1320-1323.	1.8	2
9	Complete Genome Sequences of Bordetella pertussis Clinical Isolate FR5810 and Reference Strain Tohama from Combined Oxford Nanopore and Illumina Sequencing. Microbiology Resource Announcements, 2018, 7, .	0.6	11
10	Genomic Sequencing of <i>Bordetella pertussis</i> for Epidemiology and Global Surveillance of Whooping Cough. Emerging Infectious Diseases, 2018, 24, 988-994.	4.3	29
11	Characterization of Post-Translational Modifications and Cytotoxic Properties of the Adenylate-Cyclase Hemolysin Produced by Various Bordetella pertussis and Bordetella parapertussis Isolates. Toxins, 2017, 9, 304.	3.4	6
12	Bordetella holmesii: Lipid A Structures and Corresponding Genomic Sequences Comparison in Three Clinical Isolates and the Reference Strain ATCC 51541. International Journal of Molecular Sciences, 2017, 18, 1080.	4.1	6
13	The length of poly(C) stretch in the Bordetella pertussis Pfim3 promoter determines the vag or vrg function of the fim3 gene. Microbiology (United Kingdom), 2017, 163, 1364-1368.	1.8	1
14	Structure activity characterization of Bordetella petrii lipid A, from environment to human isolates. Biochimie, 2016, 120, 87-95.	2.6	6
15	New Data on Vaccine Antigen Deficient Bordetella pertussis Isolates. Vaccines, 2015, 3, 751-770.	4.4	43
16	<i>Bordetella pertussis</i> , <i>B. parapertussis</i> , vaccines and cycles of whooping cough. Pathogens and Disease, 2015, 73, ftv055.	2.0	32
17	Global Population Structure and Evolution of Bordetella pertussis and Their Relationship with Vaccination. MBio, 2014, 5, e01074.	4.1	257
18	<i>Bordetella holmesii</i> : Comparison of Two Isolates from Blood and a Respiratory Sample. Advances in Infectious Diseases, 2013, 03, 123-133.	0.2	8

Valérie Bouchez

#	Article	IF	CITATION
19	Is the Sequenced <i>Bordetella pertussis</i> Strain Tohama I Representative of the Species?. Journal of Clinical Microbiology, 2008, 46, 2125-2128.	3.9	39
20	Genomic Content of Bordetella pertussis Clinical Isolates Circulating in Areas of Intensive Children Vaccination. PLoS ONE, 2008, 3, e2437.	2.5	57
21	Pertussis in Argentina and France. Vaccine, 2007, 25, 4335-4339.	3.8	19