Valérie Bouchez

List of Publications by Year in descending order

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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 Population Structure and Evolution of Bordetella pertussis and Their Relationship with Vaccination. MBio, 2014, 5, e01074.	4.1	257
2	Genomic Content of Bordetella pertussis Clinical Isolates Circulating in Areas of Intensive Children Vaccination. PLoS ONE, 2008, 3, e2437.	2.5	57
3	Corynebacterium rouxii sp. nov., a novel member of the diphtheriae species complex. Research in Microbiology, 2020, 171, 122-127.	2.1	45
4	New Data on Vaccine Antigen Deficient Bordetella pertussis Isolates. Vaccines, 2015, 3, 751-770.	4.4	43
5	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
6	<i>Bordetella pertussis</i> , <i>B. parapertussis</i> , vaccines and cycles of whooping cough. Pathogens and Disease, 2015, 73, ftv055.	2.0	32
7	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
8	Ongoing diphtheria outbreak in Yemen: a cross-sectional and genomic epidemiology study. Lancet Microbe, The, 2021, 2, e386-e396.	7.3	26
9	Global spatial dynamics and vaccine-induced fitness changes of <i>Bordetella pertussis</i> Science Translational Medicine, 2022, 14, eabn3253.	12.4	22
10	Pertussis in Argentina and France. Vaccine, 2007, 25, 4335-4339.	3.8	19
11	Evolution of Bordetella pertussis over a 23-year period in France, 1996 to 2018. Eurosurveillance, 2021, 26, .	7.0	15
11		7.0	15
	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,		
12	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. Complete Genome Sequences of Bordetella pertussis Clinical Isolate FR5810 and Reference Strain Tohama from Combined Oxford Nanopore and Illumina Sequencing. Microbiology Resource	1.8	14
12	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. 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, . A comprehensive resource for Bordetella genomic epidemiology and biodiversity studies. Nature	0.6	14
12 13 14	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. 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, . A comprehensive resource for Bordetella genomic epidemiology and biodiversity studies. Nature Communications, 2022, 13, . <i> Bordetella parapertussis < / i> Bacteremia: Clinical Expression and Bacterial Genomics. Open Forum</i>	1.8 0.6 12.8	14 11 11
12 13 14	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. 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, . A comprehensive resource for Bordetella genomic epidemiology and biodiversity studies. Nature Communications, 2022, 13, . 		

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19	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
20	Genome characteristics of Bordetella pertussis isolates from Tunisia. Journal of Medical Microbiology, 2019, 68, 1320-1323.	1.8	2
21	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