Astrid Vabret

List of Publications by Year
in descending order

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50 papers	2,726 citations	201658 27 h-index	182417 51 g-index
55 all docs	55 docs citations	55 times ranked	3483 citing authors

#	Article	IF	Citations
1	Molecular epidemiology of Enteroviruses and Rhinoviruses in patients with acute respiratory infections in Yaounde, Cameroon. Influenza and Other Respiratory Viruses, 2021, 15, 641-650.	3.4	8
2	Phylogenetic variability of Human Metapneumovirus in patients with acute respiratory infections in Cameroon, 2011â€“2014. Journal of Infection and Public Health, 2020, 13, 606-612.	4.1	4
3	When a viral eruption hides another one: intrafamilial outbreak of parvovirus B19 and measles virus co-infections: case report. BMC Infectious Diseases, 2020, 20, 496.	2.9	2
4	Fatal Measles Inclusion-Body Encephalitis in Adult with Untreated AIDS, France. Emerging Infectious Diseases, 2020, 26, 2231-2234.	4.3	21
5	True Measles Cases Undetected by Reverse Transcription-PCR (RT-PCR): Effect of Genetic Variability on Assay Sensitivity Needs To Be Regularly Surveyed. Journal of Clinical Microbiology, 2019, 57, .	3.9	5
6	Effectiveness of an intervention campaign on influenza vaccination of professionals in nursing homes: A cluster-randomized controlled trial. Vaccine, 2019, 37, 1260-1265.	3.8	16
7	Measles Transmission in a Fully Vaccinated Closed Cohort. Pediatric Infectious Disease Journal, 2019, 38, e230-e232.	2.0	6
8	SARS-CoV related Betacoronavirus and diverse Alphacoronavirus members found in western old-world. Virology, 2018, 517, 88-97.	2.4	71
9	Genetic diversity of human respiratory syncytial virus isolated among children with acute respiratory infections in Southern Cameroon during three consecutive epidemic seasons, 2011â€“2013. Tropical Medicine and Health, 2018, 46, 7.	2.8	12
10	Molecular characterization of human adenovirus associated with acute respiratory infections in Cameroon from 2011 to 2014. Virology Journal, 2018, 15, 153.	3.4	13
11	Contemporaneous data on the prevalence of Human Respiratory Syncytial Virus infection in people with acute respiratory tract infections in Africa (2000â€“2017). Data in Brief, 2018, 20, 940-947.	1.0	0
12	p53 regulates CD46 expression and measles virus infection in myeloma cells. Blood Advances, 2018, 2, 3492-3505.	5.2	17
13	Prevalence of human respiratory syncytial virus infection in people with acute respiratory tract infections in Africa: A systematic review and metaâ€analysis. Influenza and Other Respiratory Viruses, 2018, 12, 793-803.	3.4	18
14	Complete Genome Sequence of a Wild-Type Measles Virus Isolated during a 2016 Winter Outbreak in a Refugee Settlement in Calais, France. Genome Announcements, 2017, 5, .	0.8	2
15	First Complete Genome Sequence of a French Bovine coronavirus Strain. Genome Announcements, 2017, 5, .	0.8	0
16	Phylogenic analysis of human bocavirus detected in children with acute respiratory infection in Yaounde, Cameroon. BMC Research Notes, 2017, 10, 293.	1.4	13
17	Performance Evaluation of the VIDASÂ® Measles IgG Assay and Its Diagnostic Value for Measuring IgG Antibody Avidity in Measles Virus Infection. Viruses, 2016, 8, 234.	3.3	4
18	Comparative molecular epidemiology of two closely related coronaviruses, bovine coronavirus (BCoV) and human coronavirus OC43 (HCoV-OC43), reveals a different evolutionary pattern. Infection, Genetics and Evolution, 2016, 40, 186-191.	2.3	38

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19	Viral etiology of severe acute respiratory infections in hospitalized children in Cameroon, 2011â€“2013. Influenza and Other Respiratory Viruses, 2016, 10, 386-393.	3.4	61
20	Measles inclusion-body encephalitis (MIBE) in a immunocompromised patient. Journal of Clinical Virology, 2016, 81, 43-46.	3.1	15
21	Evaluation of Four Commercial Multiplex Molecular Tests for the Diagnosis of Acute Respiratory Infections. PLoS ONE, 2015, 10, e0130378.	2.5	58
22	Genomic Analysis of 15 Human Coronaviruses OC43 (HCoV-OC43s) Circulating in France from 2001 to 2013 Reveals a High Intra-Specific Diversity with New Recombinant Genotypes. Viruses, 2015, 7, 2358-2377.	3.3	71
23	Viral Etiology of Respiratory Tract Infections in Children at the Pediatric Hospital in Ouagadougou (Burkina Faso). PLoS ONE, 2014, 9, e110435.	2.5	44
24	Development of an efficient qRT-PCR assay for quality control and cellular quantification of respiratory samples. Journal of Clinical Virology, 2014, 60, 270-275.	3.1	16
25	First detection of equine coronavirus (ECoV) in Europe. Veterinary Microbiology, 2014, 171, 206-209.	1.9	48
26	Rapid molecular diagnosis of measles virus infection in an epidemic setting. Journal of Medical Virology, 2013, 85, 723-730.	5.0	19
27	Comparative Evaluation of Six Commercialized Multiplex PCR Kits for the Diagnosis of Respiratory Infections. PLoS ONE, 2013, 8, e72174.	2.5	75
28	Viral Etiology of Influenza-Like Illnesses in Cameroon, Januaryâ€“December 2009. Journal of Infectious Diseases, 2012, 206, S29-S35.	4.0	71
29	Fatal Measles without Rash in Immunocompetent Adult, France. Emerging Infectious Diseases, 2012, 18, 521-523.	4.3	9
30	Epidemiological and phylogenic study of human metapneumovirus infections during three consecutive outbreaks in Normandy, France. Journal of Medical Virology, 2011, 83, 517-524.	5.0	16
31	Viral etiology of respiratory infections in children under 5 years old living in tropical rural areas of Senegal: The EVIRA project. Journal of Medical Virology, 2010, 82, 866-872.	5.0	64
32	Use of a multiplex PCR/RTâ€PCR approach to assess the viral causes of influenzaâ€like illnesses in Cambodia during three consecutive dry seasons. Journal of Medical Virology, 2010, 82, 1762-1772.	5.0	36
33	Culturing the Unculturable: Human Coronavirus HKU1 Infects, Replicates, and Produces Progeny Virions in Human Ciliated Airway Epithelial Cell Cultures. Journal of Virology, 2010, 84, 11255-11263.	3.4	120
34	Burden of disease due to human coronavirus NL63 infections and periodicity of infection. Journal of Clinical Virology, 2010, 48, 104-108.	3.1	33
35	Simultaneous detection of respiratory viruses in children with acute respiratory infection using two different multiplex reverse transcription-PCR assays. Journal of Virological Methods, 2009, 162, 40-45.	2.1	41
36	Human (nonâ€severe acute respiratory syndrome) coronavirus infections in hospitalised children in France. Journal of Paediatrics and Child Health, 2008, 44, 176-181.	0.8	93

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37	Baculovirus expression of HCoV-OC43 nucleocapsid protein and development of a Western blot assay for detection of human antibodies against HCoV-OC43. Journal of Virological Methods, 2007, 139, 175-180.	2.1	20
38	Detection of the New Human Coronavirus HKU1: A Report of 6 Cases. Clinical Infectious Diseases, 2006, 42, 634-639.	5.8	172
39	Comparison of multiplex PCR assays and conventional techniques for the diagnostic of respiratory virus infections in children admitted to hospital with an acute respiratory illness. Journal of Medical Virology, 2006, 78, 1498-1504.	5.0	173
40	Inter- and intra-variant genetic heterogeneity of human coronavirus OC43 strains in France. Journal of General Virology, 2006, 87, 3349-3353.	2.9	19
41	Replication of respiratory viruses, particularly influenza virus, rhinovirus, and coronavirus in HuH7 hepatocarcinoma cell line. Journal of Medical Virology, 2005, 77, 295-301.	5.0	43
42	Human Coronavirus NL63, France. Emerging Infectious Diseases, 2005, 11, 1225-1229.	4.3	204
43	Human Respiratory Coronavirus OC43: Genetic Stability and Neuroinvasion. Journal of Virology, 2004, 78, 8824-8834.	3.4	150
44	Development of a PCR-and hybridization-based assay (PCR Adenovirus ConsensusÂ®) for the detection and the species identification of adenoviruses in respiratory specimens. Journal of Clinical Virology, 2004, 31, 116-122.	3.1	43
45	An Outbreak of Coronavirus OC43 Respiratory Infection in Normandy, France. Clinical Infectious Diseases, 2003, 36, 985-989.	5.8	210
46	Coronavirus 229E-Related Pneumonia in Immunocompromised Patients. Clinical Infectious Diseases, 2003, 37, 929-932.	5.8	253
47	Direct diagnosis of human respiratory coronaviruses 229E and OC43 by the polymerase chain reaction. Journal of Virological Methods, 2001, 97, 59-66.	2.1	79
48	Comparison of three non-nested RT-PCR for the detection of influenza A viruses. Journal of Clinical Virology, 2000, 17, 167-175.	3.1	29
49	Detection of Respiratory Syncytial Virus A and B and Parainfluenzavirus 3 Sequences in Respiratory Tracts of Infants by a Single PCR with Primers Targeted to the L-Polymerase Gene and Differential Hybridization. Journal of Clinical Microbiology, 1998, 36, 796-801.	3.9	59
50	Detection of respiratory syncytial virus, parainfluenzavirus 3, adenovirus and rhinovirus sequences in respiratory tract of infants by polymerase chain reaction and hybridization. Clinical and Diagnostic Virology, 1997, 8, 31-40.	1.7	118