Guillaume Croville

List of Publications by Year
in descending order

20 papers	438 citations	777949 13 h-index	889612 19 g-index
22 all docs	22 docs citations	22 times ranked	820 citing authors

#	Article	IF	Citations
1	First Detection and Identification of FAdV-8b as the Causative Agent of an Outbreak of Inclusion Body Hepatitis in a Commercial Broiler Farm in Greece. Veterinary Sciences, 2022, 9, 160.	0.6	0
2	Highly Pathogenic Avian Influenza A(H5N8) Clade 2.3.4.4b Virus in Dust Samples from Poultry Farms, France, 2021. Emerging Infectious Diseases, 2022, 28, 1446-1450.	2.0	20
3	Viral tropism and detection of clade 2.3.4.4b H5N8 highly pathogenic avian influenza viruses in feathers of ducks and geese. Scientific Reports, 2021, 11, 5928.	1.6	15
4	Household Cases Suggest That Cats Belonging to Owners with COVID-19 Have a Limited Role in Virus Transmission. Viruses, 2021, 13, 673.	1.5	12
5	Role of Backyard Flocks in Transmission Dynamics of Highly Pathogenic Avian Influenza A(H5N8) Clade 2.3.4.4, France, 2016â€“2017. Emerging Infectious Diseases, 2019, 25, 551-554.	2.0	21
6	Disclosing respiratory co-infections: a broad-range panel assay for avian respiratory pathogens on a nanofluidic PCR platform. Avian Pathology, 2018, 47, 253-260.	0.8	12
7	Rapid whole-genome based typing and surveillance of avipoxviruses using nanopore sequencing. Journal of Virological Methods, 2018, 261, 34-39.	1.0	31
8	Intra-host viral variability in children clinically infected with H1N1 (2009) pandemic influenza. Infection, Genetics and Evolution, 2015, 33, 47-54.	1.0	8
9	Subpopulations in aMPV vaccines are unlikely to be the only cause of reversion to virulence. Vaccine, 2015, 33, 2438-2441.	1.7	15
10	Full genome sequence of guinea fowl coronavirus associated with fulminating disease. Virus Genes, 2015, 50, 514-517.	0.7	10
11	Protection Patterns in Duck and Chicken after Homo- or Hetero-Subtypic Reinfections with H5 and H7 Low Pathogenicity Avian Influenza Viruses: A Comparative Study. PLoS ONE, 2014, 9, e105189.	1.1	14
12	Complete Genome Sequence of a Field Strain of Peste des Petits Ruminants Virus Isolated during 2010-2014 Epidemics in Senegal. Genome Announcements, 2014, 2, .	0.8	14
13	Novel Avian Coronavirus and Fulminating Disease in Guinea Fowl, France. Emerging Infectious Diseases, 2014, 20, 105-8.	2.0	34
14	Deletion of the C-terminal ESEV domain of NS1 does not affect the replication of a low-pathogenic avian influenza virus H7N1 in ducks and chickens. Journal of General Virology, 2013, 94, 50-58.	1.3	27
15	Whole-genome, deep pyrosequencing analysis of a duck influenza A virus evolution in swine cells. Infection, Genetics and Evolution, 2013, 18, 31-41.	1.0	19
16	A low-pathogenic avian influenza H6N1 outbreak in a turkey flock in France: a comprehensive case report. Avian Pathology, 2012, 41, 569-577.	0.8	18
17	Length Variations in the NA Stalk of an H7N1 Influenza Virus Have Opposite Effects on Viral Excretion in Chickens and Ducks. Journal of Virology, 2012, 86, 584-588.	1.5	49
18	Field Monitoring of Avian Influenza Viruses: Whole-Genome Sequencing and Tracking of Neuraminidase Evolution Using 454 Pyrosequencing. Journal of Clinical Microbiology, 2012, 50, 2881-2887.	1.8	25

#	Article	IF	Citations
19	Actions of the prototypical 5-HT1A receptor agonist 8-OH-DPAT at human Î±2-adrenoceptors: (+)8-OH-DPAT, but not (â^')8-OH-DPAT is an Î±2B subtype preferential agonist. European Journal of Pharmacology, 2010, 640, 8-14.	1.7	4
20	Species-Specific Contribution of the Four C-Terminal Amino Acids of Influenza A Virus NS1 Protein to Virulence. Journal of Virology, 2010, 84, 6733-6747.	1.5	88