

# Mariette F Ducatez

## List of Publications by Year in descending order

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Version: 2024-02-01

55  
papers

1,711  
citations

394421

19  
h-index

289244

40  
g-index

57  
all docs

57  
docs citations

57  
times ranked

1651  
citing authors

#	ARTICLE	IF	CITATIONS
1	Influenza D virus in respiratory disease in Canadian, province of QuÃ©bec, cattle: Relative importance and evidence of new reassortment between different clades. <i>Transboundary and Emerging Diseases</i> , 2022, 69, 1227-1245.	3.0	14
2	Hydroxychloroquine inhibits proteolytic processing of endogenous TLR7 protein in human primary plasmacytoid dendritic cells. <i>European Journal of Immunology</i> , 2022, 52, 54-61.	2.9	10
3	First expert elicitation of knowledge on drivers of emergence of influenza D in Europe. <i>Transboundary and Emerging Diseases</i> , 2021, 68, 3349-3359.	3.0	9
4	Formal estimation of the seropositivity cut-off of the hemagglutination inhibition assay in field diagnosis of influenza D virus in cattle and estimation of the associated true prevalence in Morocco. <i>Transboundary and Emerging Diseases</i> , 2021, 68, 1392-1399.	3.0	5
5	Emerging Influenza D virus infection in European livestock as determined in serology studies: Are we underestimating its spread over the continent?. <i>Transboundary and Emerging Diseases</i> , 2021, 68, 1125-1135.	3.0	18
6	Antigenic and molecular characterization of low pathogenic avian influenza A(H9N2) viruses in sub-Saharan Africa from 2017 through 2019. <i>Emerging Microbes and Infections</i> , 2021, 10, 753-761.	6.5	10
7	Protective Efficacy Evaluation of Four Inactivated Commercial Vaccines Against Low Pathogenic Avian Influenza H9N2 Virus Under Experimental Conditions in Broiler Chickens. <i>Avian Diseases</i> , 2021, 65, 351-357.	1.0	8
8	Pathogenesis of Avian Influenza Virus Subtype H9N2 in Turkeys and Evaluation of Inactivated Vaccine Efficacy. <i>Avian Diseases</i> , 2021, 65, 46-51.	1.0	2
9	Serological Surveillance of Influenza D Virus in Ruminants and Swine in West and East Africa, 2017-2020. <i>Viruses</i> , 2021, 13, 1749.	3.3	11
10	Enhanced Pathogenesis Caused by Influenza D Virus and <i>Mycoplasma bovis</i> Coinfection in Calves: a Disease Severity Linked with Overexpression of IFN- $\beta$ as a Key Player of the Enhanced Innate Immune Response in Lungs. <i>Microbiology Spectrum</i> , 2021, 9, e0169021.	3.0	16
11	Co-infections of chickens with avian influenza virus H9N2 and Moroccan Italy 02 infectious bronchitis virus: effect on pathogenesis and protection conferred by different vaccination programmes. <i>Avian Pathology</i> , 2020, 49, 21-28.	2.0	22
12	Complete genome analysis and time scale evolution of very virulent infectious bursal disease viruses isolated from recent outbreaks in Morocco. <i>Infection, Genetics and Evolution</i> , 2020, 77, 104097.	2.3	9
13	Full-length genome sequences of the first H9N2 avian influenza viruses isolated in the Northeast of Algeria. <i>Virology Journal</i> , 2020, 17, 108.	3.4	12
14	CD5 signalosome coordinates antagonist TCR signals to control the generation of Treg cells induced by foreign antigens. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 12969-12979.	7.1	15
15	Risk assessment for influenza D in Europe. <i>EFSA Supporting Publications</i> , 2020, 17, 1853E.	0.7	2
16	First report of influenza D virus infection in Turkish cattle with respiratory disease. <i>Research in Veterinary Science</i> , 2020, 130, 98-102.	1.9	10
17	Risk Mapping of Influenza D Virus Occurrence in Ruminants and Swine in Togo Using a Spatial Multicriteria Decision Analysis Approach. <i>Viruses</i> , 2020, 12, 128.	3.3	16
18	Molecular epidemiology of respiratory viruses in commercial chicken flocks in Pakistan from 2014 through to 2016. <i>BMC Veterinary Research</i> , 2019, 15, 351.	1.9	14

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19	Pathogenesis, Host Innate Immune Response, and Aerosol Transmission of Influenza D Virus in Cattle. <i>Journal of Virology</i> , 2019, 93, .	3.4	48
20	Serological Evidence of Influenza D Virus Circulation Among Cattle and Small Ruminants in France. <i>Viruses</i> , 2019, 11, 516.	3.3	43
21	Seroprevalence of influenza D virus in selected sample groups of Irish cattle, sheep and pigs. <i>Irish Veterinary Journal</i> , 2019, 72, 11.	2.1	25
22	Disclosing respiratory co-infections: a broad-range panel assay for avian respiratory pathogens on a nanofluidic PCR platform. <i>Avian Pathology</i> , 2018, 47, 253-260.	2.0	12
23	Genotypic characterisation of Avian paramyxovirus type-1 viruses isolated from aquatic birds in Uganda. <i>Onderstepoort Journal of Veterinary Research</i> , 2018, 85, e1-e7.	1.2	4
24	Influenza D Virus in Cattle, Ireland. <i>Emerging Infectious Diseases</i> , 2018, 24, 389-391.	4.3	66
25	Influenza D Virus Circulation in Cattle and Swine, Luxembourg, 2012â€“2016. <i>Emerging Infectious Diseases</i> , 2018, 24, 1388-1389.	4.3	52
26	Low Pathogenic Avian Influenza and Coinfecting Pathogens: A Review of Experimental Infections in Avian Models. <i>Avian Diseases</i> , 2017, 61, 3-15.	1.0	38
27	Efficacy of Massachusetts and 793B Vaccines Against Infectious Bronchitis Moroccan-Italy O2 Virus in Specific-Pathogen-Free Chickens and Commercial Broilers. <i>Avian Diseases</i> , 2017, 61, 466-471.	1.0	5
28	Serologic Evidence for Influenza C and D Virus among Ruminants and Camelids, Africa, 1991â€“2015. <i>Emerging Infectious Diseases</i> , 2017, 23, 1556-1559.	4.3	104
29	Highly pathogenic avian influenza H5N1 clade 2.3.2.1 and clade 2.3.4 viruses do not induce a clade-specific phenotype in mallard ducks. <i>Journal of General Virology</i> , 2017, 98, 1232-1244.	2.9	10
30	First outbreaks and phylogenetic analyses of avian influenza H9N2 viruses isolated from poultry flocks in Morocco. <i>Virology Journal</i> , 2016, 13, 140.	3.4	46
31	Recommendations for a standardized avian coronavirus (AvCoV) nomenclature: outcome from discussions within the framework of the European Union COST Action FA1207: â€œtowards control of avian coronaviruses: strategies for vaccination, diagnosis and surveillanceâ€*. <i>Avian Pathology</i> , 2016, 45, 602-603.	2.0	6
32	Wholeâ€“genome analysis of influenza A(H1N1)pdm09 viruses isolated in Uganda from 2009 to 2011. <i>Influenza and Other Respiratory Viruses</i> , 2016, 10, 486-492.	3.4	11
33	Low pathogenic avian influenza (H9N2) in chicken: Evaluation of an ancestral H9-MVA vaccine. <i>Veterinary Microbiology</i> , 2016, 189, 59-67.	1.9	11
34	Pandemic Seasonal H1N1 Reassortants Recovered from Patient Material Display a Phenotype Similar to That of the Seasonal Parent. <i>Journal of Virology</i> , 2016, 90, 7647-7656.	3.4	0
35	Phylogenetic analysis of avian infectious bronchitis virus S1 glycoprotein regions reveals emergence of a new genotype in Moroccan broiler chicken flocks. <i>Virology Journal</i> , 2015, 12, 116.	3.4	14
36	Changes to the dynamic nature of hemagglutinin and the emergence of the 2009 pandemic H1N1 influenza virus. <i>Scientific Reports</i> , 2015, 5, 12828.	3.3	10

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37	Influenza D Virus in Cattle, France, 2011â€“2014. <i>Emerging Infectious Diseases</i> , 2015, 21, 368-71.	4.3	134
38	Influenza A(H1N1)pdm09 virus in pigs, Togo, 2013. <i>Veterinary Microbiology</i> , 2015, 177, 201-205.	1.9	16
39	Full genome sequence of guinea fowl coronavirus associated with fulminating disease. <i>Virus Genes</i> , 2015, 50, 514-517.	1.6	10
40	Time scale evolution of avipoxviruses. <i>Infection, Genetics and Evolution</i> , 2015, 35, 75-81.	2.3	6
41	Identification of a Novel Coronavirus from Guinea Fowl Using Metagenomics. <i>Methods in Molecular Biology</i> , 2015, 1282, 27-31.	0.9	6
42	Novel Avian Coronavirus and Fulminating Disease in Guinea Fowl, France. <i>Emerging Infectious Diseases</i> , 2014, 20, 105-8.	4.3	34
43	High pathogenicity and low genetic evolution of avian paramyxovirus type I (Newcastle disease virus) isolated from live bird markets in Uganda. <i>Virology Journal</i> , 2014, 11, 173.	3.4	30
44	Diversity of avipoxviruses in captive-bred Houbara bustard. <i>Veterinary Research</i> , 2014, 45, 98.	3.0	12
45	Prevalence of influenza A viruses in livestock and free-living waterfowl in Uganda. <i>BMC Veterinary Research</i> , 2014, 10, 50.	1.9	18
46	Long-term vaccine-induced heterologous protection against H5N1 influenza viruses in the ferret model. <i>Influenza and Other Respiratory Viruses</i> , 2013, 7, 506-512.	3.4	6
47	Isolation of a Novel Swine Influenza Virus from Oklahoma in 2011 Which Is Distantly Related to Human Influenza C Viruses. <i>PLoS Pathogens</i> , 2013, 9, e1003176.	4.7	268
48	Active Surveillance for Influenza A Virus among Swine, Midwestern United States, 2009â€“2011. <i>Emerging Infectious Diseases</i> , 2013, 19, 954-960.	4.3	66
49	A low-pathogenic avian influenza H6N1 outbreak in a turkey flock in France: a comprehensive case report. <i>Avian Pathology</i> , 2012, 41, 569-577.	2.0	18
50	Both influenza hemagglutinin and polymerase acidic genes are important for delayed pandemic 2009 H1N1 virus clearance in the ferret model. <i>Virology</i> , 2012, 432, 389-393.	2.4	6
51	Surveillance for Influenza Viruses in Poultry and Swine, West Africa, 2006â€“2008. <i>Emerging Infectious Diseases</i> , 2012, 18, 1446-1452.	4.3	37
52	Feasibility of reconstructed ancestral H5N1 influenza viruses for cross-clade protective vaccine development. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 349-354.	7.1	52
53	Extent of Antigenic Cross-Reactivity among Highly Pathogenic H5N1 Influenza Viruses. <i>Journal of Clinical Microbiology</i> , 2011, 49, 3531-3536.	3.9	28
54	Multiple Reassortment between Pandemic (H1N1) 2009 and Endemic Influenza Viruses in Pigs, United States. <i>Emerging Infectious Diseases</i> , 2011, 17, 1624-1629.	4.3	165

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55	Newcastle disease virus in West Africa: new virulent strains identified in non-commercial farms. Archives of Virology, 2009, 154, 47-54.	2.1	77