

Influenza A Virus Migration and Persistence in North A

PLoS Pathogens

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Genetic Structure of Avian Influenza Viruses from Ducks of the Atlantic Flyway of North America. PLoS ONE, 2014, 9, e86999.	2.5	14
2	Potential Role of Migratory Quail in Spreading of Some Zoonotic Pathogens in Egypt. American Journal of Animal and Veterinary Sciences, 2014, 9, 203-210.	0.5	5
3	Cross-conservation of T-cell epitopes. Human Vaccines and Immunotherapeutics, 2014, 10, 256-262.	3.3	22
4	Bird harvesting practices and knowledge, risk perceptions, and attitudes regarding avian influenza among Canadian First Nations subsistence hunters: implications for influenza pandemic plans. BMC Public Health, 2014, 14, 1113.	2.9	6
5	Adaptive Evolution and Environmental Durability Jointly Structure Phylodynamic Patterns in Avian Influenza Viruses. PLoS Biology, 2014, 12, e1001931.	5.6	36
6	Using quantitative disease dynamics as a tool for guiding response to avian influenza in poultry in the United States of America. Preventive Veterinary Medicine, 2014, 113, 376-397.	1.9	19
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8	Avian Influenza from an Ecohealth Perspective. EcoHealth, 2014, 11, 4-14.	2.0	13
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11	Global Avian Influenza Surveillance in Wild Birds: A Strategy to Capture Viral Diversity. Emerging Infectious Diseases, 2015, 21, e1-7.	4.3	46
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16	Spread and Persistence of Influenza A Viruses in Waterfowl Hosts in the North American Mississippi Migratory Flyway. Journal of Virology, 2015, 89, 5371-5381.	3.4	29
17	Influenza A virus evolution and spatio-temporal dynamics in Eurasian wild birds: a phylogenetic and phylogeographical study of whole-genome sequence data. Journal of General Virology, 2015, 96, 2050-2060.	2.9	23
18	The Genetic Diversity of Influenza A Viruses in Wild Birds in Peru. PLoS ONE, 2016, 11, e0146059.	2.5	24

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19	Ecosystem Interactions Underlie the Spread of Avian Influenza A Viruses with Pandemic Potential. <i>PLoS Pathogens</i> , 2016, 12, e1005620.	4.7	48
20	Identification of migratory bird flyways in North America using community detection on biological networks. <i>Ecological Applications</i> , 2016, 26, 740-751.	3.8	27
21	Extensive Allelic Diversity of MHC Class I in Wild Mallard Ducks. <i>Journal of Immunology</i> , 2016, 197, 783-794.	0.8	14
22	Transmission of influenza reflects seasonality of wild birds across the annual cycle. <i>Ecology Letters</i> , 2016, 19, 915-925.	6.4	59
23	A point mutation in the polymerase protein PB2 allows a reassortant H9N2 influenza isolate of wild-bird origin to replicate in human cells. <i>Infection, Genetics and Evolution</i> , 2016, 41, 279-288.	2.3	4
24	Genome-scale phylodynamics and evolution analysis of global H7N7 influenza viruses. <i>Veterinary Microbiology</i> , 2016, 193, 83-92.	1.9	2
25	Analysis of influenza A viruses from gulls: An evaluation of inter-regional movements and interactions with other avian and mammalian influenza A viruses. <i>Cogent Biology</i> , 2016, 2, 1234957.	1.7	9
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27	Inference of Japanese encephalitis virus ecological and evolutionary dynamics from passive and active virus surveillance. <i>Virus Evolution</i> , 2016, 2, vew009.	4.9	6
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30	THE PATHOGENESIS OF CLADE 2.3.4.4 H5 HIGHLY PATHOGENIC AVIAN INFLUENZA VIRUSES IN RUDDY DUCK (<i>Oxyura jamaicensis</i>) AND LESSER SCAUP (<i>Aythya affinis</i>). <i>Journal of Wildlife Diseases</i> , 2017, 53, 832-842.	0.8	20
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39	A Review of Eight High-Priority, Economically Important Viral Pathogens of Poultry within the Caribbean Region. <i>Veterinary Sciences</i> , 2018, 5, 14.	1.7	21
40	Contrasting selective patterns across the segmented genome of bluetongue virus in a global reassortment hotspot. <i>Virus Evolution</i> , 2019, 5, vez027.	4.9	17
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50	Establishment and application of the National Parasitic Resource Center (NPRC) in China. <i>Advances in Parasitology</i> , 2020, 110, 373-400.	3.2	0
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