

Influenza in Migratory Birds and Evidence of Limited In

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Increased detection of influenza A H16 in the United States. Archives of Virology, 2008, 153, 1981-1983.	2.1	13
2	Assessing the exposure of German and Austrian bird ringers to West Nile virus (Flavivirus) and evaluating their potential risk of infection. Journal of Ornithology, 2008, 149, 271-275.	1.1	12
3	Gene Segment Reassortment Between American and Asian Lineages of Avian Influenza Virus from Waterfowl in the Beringia Area. Vector-Borne and Zoonotic Diseases, 2008, 8, 783-790.	1.5	54
4	Prevalence of Influenza A viruses in wild migratory birds in Alaska: Patterns of variation in detection at a crossroads of intercontinental flyways. Virology Journal, 2008, 5, 71.	3.4	122
5	PERPETUATION OF AVIAN INFLUENZA IN THE AMERICAS: EXAMINING THE ROLE OF SHOREBIRDS IN PATAGONIA. Auk, 2008, 125, 494-495.	1.4	6
6	Phylogenetic Diversity and Molecular Detection of Bacteria in Gull Feces. Applied and Environmental Microbiology, 2008, 74, 3969-3976.	3.1	163
7	The Evolutionary Genetics and Emergence of Avian Influenza Viruses in Wild Birds. PLoS Pathogens, 2008, 4, e1000076.	4.7	334
8	Animal health and welfare aspects of avian influenza and the risk of its introduction into the EU poultry holdings - Scientific opinion of the Panel on Animal Health and Welfare. EFSA Journal, 2008, 6, 715.	1.8	7
9	Maternal Antibody Transfer in Yellow-legged Gulls. Emerging Infectious Diseases, 2009, 15, 1147-1149.	4.3	19
10	H5N1 Surveillance in Migratory Birds in Java, Indonesia. Vector-Borne and Zoonotic Diseases, 2009, 9, 695-702.	1.5	8
11	Invasions by Eurasian Avian Influenza Virus H6 Genes and Replacement of Its North American Clade. Emerging Infectious Diseases, 2009, 15, 1040-1045.	4.3	34
12	Highly Pathogenic Avian Influenza Virus A (H7N3) in Domestic Poultry, Saskatchewan, Canada, 2007. Emerging Infectious Diseases, 2009, 15, 1492-1495.	4.3	72
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15	Gene flow and competitive exclusion of avian influenza A virus in natural reservoir hosts. Virology, 2009, 390, 289-297.	2.4	108
16	Avian influenza at both ends of a migratory flyway: characterizing viral genomic diversity to optimize surveillance plans for North America. Evolutionary Applications, 2009, 2, 457-468.	3.1	61
17	Surveillance for highly pathogenic avian influenza in wild birds in the USA. Integrative Zoology, 2009, 4, 426-439.	2.6	56
18	Genetic characterization of two low pathogenic avian influenza virus H5N1 isolates from Ontario, Canada. Virus Genes, 2009, 38, 149-154.	1.6	6

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19	Molecular character of influenza A/H1N1 2009: Implications for spread and control. <i>Indian Journal of Microbiology</i> , 2009, 49, 339-347.	2.7	5
20	Ducks: The "Trojan Horses" of H5N1 influenza. <i>Influenza and Other Respiratory Viruses</i> , 2009, 3, 121-128.	3.4	206
21	Australian surveillance for avian influenza viruses in wild birds between July 2005 and June 2007. <i>Australian Veterinary Journal</i> , 2009, 87, 266-272.	1.1	45
22	Evolutionary biology, community ecology and avian influenza research. <i>Infection, Genetics and Evolution</i> , 2009, 9, 298-303.	2.3	21
23	Companion Animals as Sentinels for Public Health. <i>Veterinary Clinics of North America - Small Animal Practice</i> , 2009, 39, 241-250.	1.5	63
24	Avian influenza virus: Of virus and bird ecology. <i>Vaccine</i> , 2009, 27, 6340-6344.	3.8	93
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27	Comparison of Cloacal and Oropharyngeal Samples for the Detection of Avian Influenza Virus in Wild Birds. <i>Avian Diseases</i> , 2010, 54, 115-119.	1.0	15
28	Hitchhiking and the Population Genetic Structure of Avian Influenza Virus. <i>Journal of Molecular Evolution</i> , 2010, 70, 98-105.	1.8	39
29	Gene segment reassortment between Eurasian and American clades of avian influenza virus in Italy. <i>Archives of Virology</i> , 2010, 155, 77-81.	2.1	9
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50	Use of phylogenetics in the molecular epidemiology and evolutionary studies of viral infections. <i>Critical Reviews in Clinical Laboratory Sciences</i> , 2010, 47, 5-49.	6.1	56
51	Limited evidence of trans-hemispheric movement of avian influenza viruses among contemporary North American shorebird isolates. <i>Virus Research</i> , 2010, 148, 44-50.	2.2	36
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53	Avian Influenza Virus Surveillance and Wild Birds: Past and Present. <i>Avian Diseases</i> , 2010, 54, 394-398.	1.0	60
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108	Natural history of highly pathogenic avian influenza H5N1. <i>Virus Research</i> , 2013, 178, 63-77.	2.2	122
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118	Development and Pre-Clinical Evaluation of Two LAIV Strains against Potentially Pandemic H2N2 Influenza Virus. <i>PLoS ONE</i> , 2014, 9, e102339.	2.5	25
119	Bird harvesting practices and knowledge, risk perceptions, and attitudes regarding avian influenza among Canadian First Nations subsistence hunters: implications for influenza pandemic plans. <i>BMC Public Health</i> , 2014, 14, 1113.	2.9	6
120	Prevalence and control of H7 avian influenza viruses in birds and humans. <i>Epidemiology and Infection</i> , 2014, 142, 896-920.	2.1	56
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125	Diverse inter-continental and host lineage reassortant avian influenza A viruses in pelagic seabirds. <i>Infection, Genetics and Evolution</i> , 2014, 22, 103-111.	2.3	15
126	Host adaptation and transmission of influenza A viruses in mammals. <i>Emerging Microbes and Infections</i> , 2014, 3, 1-10.	6.5	132
127	Determinants of virulence of influenza A virus. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2014, 33, 479-490.	2.9	77
128	Global geno-proteomic analysis reveals cross-continental sequence conservation and druggable sites among influenza virus polymerases. <i>Antiviral Research</i> , 2014, 112, 120-131.	4.1	5
129	Genetic characterization of highly pathogenic H5N1 avian influenza virus from live migratory birds in Bangladesh. <i>Virus Genes</i> , 2014, 49, 438-448.	1.6	14

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130	Influenza A Virus Reassortment. <i>Current Topics in Microbiology and Immunology</i> , 2014, 385, 377-401.	1.1	110
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143	AS03-adjuvanted H7N1 detergent-split virion vaccine is highly immunogenic in unprimed mice and induces cross-reactive antibodies to emerged H7N9 and additional H7 subtypes. <i>Vaccine</i> , 2015, 33, 3784-3787.	3.8	9
144	Epidemiology, ecology and gene pool of influenza A virus in Egypt: Will Egypt be the epicentre of the next influenza pandemic?. <i>Virulence</i> , 2015, 6, 6-18.	4.4	31
145	Genetic characterization of a rare H12N3 avian influenza virus isolated from a green-winged teal in Japan. <i>Virus Genes</i> , 2015, 50, 316-320.	1.6	6
146	Long-term surveillance of H7 influenza viruses in American wild aquatic birds: are the H7N3 influenza viruses in wild birds the precursors of highly pathogenic strains in domestic poultry?. <i>Emerging Microbes and Infections</i> , 2015, 4, 1-9.	6.5	25
147	Possible basis for the emergence of H1N1 viruses with pandemic potential from avian hosts. <i>Emerging Microbes and Infections</i> , 2015, 4, 1-10.	6.5	14

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149	Replication of live attenuated cold-adapted H2N2 influenza virus vaccine candidates in non human primates. <i>Vaccine</i> , 2015, 33, 193-200.	3.8	7
150	Extensive Allelic Diversity of MHC Class I in Wild Mallard Ducks. <i>Journal of Immunology</i> , 2016, 197, 783-794.	0.8	14
151	Limited evidence of intercontinental dispersal of avian paramyxovirus serotype 4 by migratory birds. <i>Infection, Genetics and Evolution</i> , 2016, 40, 104-108.	2.3	13
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153	Influenza A Viruses from Overwintering and Spring-Migrating Waterfowl in the Lake Erie Basin, United States. <i>Avian Diseases</i> , 2016, 60, 241-244.	1.0	4
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