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List of Publications by Year in descending order

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66911 87888 6,388 94 38 78 citations h-index g-index papers 99 99 99 6187 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	GATA6 regulates HNF4 and is required for differentiation of visceral endoderm in the mouse embryo. Genes and Development, 1998, 12, 3579-3590.	5.9	589
2	GATA-6: A Zinc Finger Transcription Factor That Is Expressed in Multiple Cell Lineages Derived from Lateral Mesoderm. Developmental Biology, 1996, 177, 309-322.	2.0	427
3	Role for migratory wild birds in the global spread of avian influenza H5N8. Science, 2016, 354, 213-217.	12.6	362
4	Myocardin Is a Critical Serum Response Factor Cofactor in the Transcriptional Program Regulating Smooth Muscle Cell Differentiation. Molecular and Cellular Biology, 2003, 23, 2425-2437.	2.3	325
5	Intercontinental Spread of Asian-Origin H5N8 to North America through Beringia by Migratory Birds. Journal of Virology, 2015, 89, 6521-6524.	3.4	306
6	Structure and Expression of a Smooth Muscle Cell-specific Gene, SM22α. Journal of Biological Chemistry, 1995, 270, 13460-13469.	3.4	240
7	GATA-5: A Transcriptional Activator Expressed in a Novel Temporally and Spatially-Restricted Pattern during Embryonic Development. Developmental Biology, 1997, 183, 21-36.	2.0	234
8	A Serum Response Factor-Dependent Transcriptional Regulatory Program Identifies Distinct Smooth Muscle Cell Sublineages. Molecular and Cellular Biology, 1997, 17, 2266-2278.	2.3	203
9	Novel Eurasian Highly Pathogenic Avian Influenza A H5 Viruses in Wild Birds, Washington, USA, 2014. Emerging Infectious Diseases, 2015, 21, 886-890.	4.3	196
10	Coordinate transcription of variant surface glycoprotein genes and an expression site associated gene family in Trypanosoma brucei. Cell, 1985, 42, 173-182.	28.9	181
11	Emergence of Fatal Avian Influenza in New England Harbor Seals. MBio, 2012, 3, e00166-12.	4.1	161
12	Evolutionary dynamics of Newcastle disease virus. Virology, 2009, 391, 64-72.	2.4	145
13	Genetic evidence of intercontinental movement of avian influenza in a migratory bird: the northern pintail (<i>Anas acuta</i>). Molecular Ecology, 2008, 17, 4754-4762.	3.9	135
14	Highly Pathogenic Avian Influenza Viruses and Generation of Novel Reassortants, United States, 2014–2015. Emerging Infectious Diseases, 2016, 22, 1283-1285.	4.3	132
15	Possibility for reverse zoonotic transmission of SARS-CoV-2 to free-ranging wildlife: A case study of bats. PLoS Pathogens, 2020, 16, e1008758.	4.7	127
16	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
17	GATA-4 Activates Transcription Via Two Novel Domains That Are Conserved within the GATA-4/5/6 Subfamily. Journal of Biological Chemistry, 1997, 272, 8515-8524.	3.4	120
18	Worldwide Phylogenetic Relationship of Avian Poxviruses. Journal of Virology, 2013, 87, 4938-4951.	3.4	112

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19	Developmental Pattern of Expression and Genomic Organization of the Calponin-h1 Gene. Journal of Biological Chemistry, 1996, 271, 395-403.	3.4	107
20	The Ets protein Spi-B is expressed exclusively in B cells and T cells during development Journal of Experimental Medicine, 1996, 184, 203-214.	8.5	101
21	Migratory flyway and geographical distance are barriers to the gene flow of influenza virus among North American birds. Ecology Letters, 2012, 15, 24-33.	6.4	86
22	SARS-CoV-2 Exposure in Escaped Mink, Utah, USA. Emerging Infectious Diseases, 2021, 27, 988-990.	4.3	78
23	Cyclic Avian Mass Mortality in the Northeastern United States Is Associated with a Novel Orthomyxovirus. Journal of Virology, 2015, 89, 1389-1403.	3.4	68
24	An invasive cleavage assay for direct quantitation of specific RNAs. Nature Biotechnology, 2001, 19, 673-676.	17.5	66
25	North Atlantic Migratory Bird Flyways Provide Routes for Intercontinental Movement of Avian Influenza Viruses. PLoS ONE, 2014, 9, e92075.	2.5	65
26	Comparison of Filters for Concentrating Microbial Indicators and Pathogens in Lake Water Samples. Applied and Environmental Microbiology, 2013, 79, 1342-1352.	3.1	63
27	Intercontinental reassortment and genomic variation of low pathogenic avian influenza viruses isolated from northern pintails (Anas acuta) in Alaska: Examining the evidence through space and time. Virology, 2010, 401, 179-189.	2.4	62
28	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
29	Transmission and reassortment of avian influenza viruses at the Asian–North American interface. Virology, 2010, 406, 352-359.	2.4	55
30	Characterization of Low-Pathogenicity H5N1 Avian Influenza Viruses from North America. Journal of Virology, 2007, 81, 11612-11619.	3.4	54
31	Respiratory transmission of an avian H3N8 influenza virus isolated from a harbour seal. Nature Communications, 2014, 5, 4791.	12.8	54
32	Experimental challenge of a North American bat species, big brown bat (<i>Eptesicus fuscus</i>), with SARS oVâ€2. Transboundary and Emerging Diseases, 2021, 68, 3443-3452.	3.0	54
33	Genetic diversity and mutation of avian paramyxovirus serotype 1 (Newcastle disease virus) in wild birds and evidence for intercontinental spread. Archives of Virology, 2013, 158, 2495-2503.	2.1	53
34	Interspecific exchange of avian influenza virus genes in Alaska: the influence of transâ€hemispheric migratory tendency and breeding ground sympatry. Molecular Ecology, 2011, 20, 1015-1025.	3.9	47
35	Influence of Body Condition on Influenza A Virus Infection in Mallard Ducks: Experimental Infection Data. PLoS ONE, 2011, 6, e22633.	2.5	46
36	Novel H5 Clade 2.3.4.4 Reassortant (H5N1) Virus from a Green-Winged Teal in Washington, USA. Genome Announcements, 2015, 3, .	0.8	45

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37	Experimental Infection of a North American Raptor, American Kestrel (Falco sparverius), with Highly Pathogenic Avian Influenza Virus (H5N1). PLoS ONE, 2009, 4, e7555.	2.5	44
38	Fluid Spatial Dynamics of West Nile Virus in the United States: Rapid Spread in a Permissive Host Environment. Journal of Virology, 2016, 90, 862-872.	3.4	42
39	Mapping of RNA accessible sites by extension of random oligonucleotide libraries with reverse transcriptase. Rna, 2001, 7, 314-327.	3.5	41
40	Analytical Validation of a Real-Time Reverse Transcription Polymerase Chain Reaction Test for Pan-American Lineage H7 Subtype Avian Influenza Viruses. Journal of Veterinary Diagnostic Investigation, 2008, 20, 612-616.	1.1	39
41	Surveillance for zoonotic and selected pathogens in harbor seals Phoca vitulina from central California. Diseases of Aquatic Organisms, 2014, 111, 93-106.	1.0	37
42	Presence of Avian Influenza Viruses in Waterfowl and Wetlands during Summer 2010 in California: Are Resident Birds a Potential Reservoir?. PLoS ONE, 2012, 7, e31471.	2.5	37
43	Limited evidence of trans-hemispheric movement of avian influenza viruses among contemporary North American shorebird isolates. Virus Research, 2010, 148, 44-50.	2.2	36
44	Discovery of a Novel Hepatovirus (<i>Phopivirus</i> of Seals) Related to Human Hepatitis A Virus. MBio, 2015, 6, .	4.1	36
45	Evolution of a reassortant North American gull influenza virus lineage: drift, shift and stability. Virology Journal, 2013, 10, 179.	3.4	34
46	Leishmania mexicana mexicana gp63 is a site-specific neutral endopeptidase. Molecular and Biochemical Parasitology, 1990, 40, 163-172.	1.1	33
47	Surveillance for High Pathogenicity Avian Influenza Virus in Wild Birds in the Pacific Flyway of the United States, 2006–2007. Avian Diseases, 2009, 53, 222-230.	1.0	33
48	Impact of West Nile virus and other mortality factors on American white pelicans at breeding colonies in the northern plains of North America. Biological Conservation, 2008, 141, 1021-1031.	4.1	32
49	Influenza A Virus Infections in Land Birds, People's Republic of China. Emerging Infectious Diseases, 2008, 14, 1644-1646.	4.3	31
50	Whole-genome sequencing of genotype VI Newcastle disease viruses from formalin-fixed paraffin-embedded tissues from wild pigeons reveals continuous evolution and previously unrecognized genetic diversity in the U.S Virology Journal, 2018, 15, 9.	3.4	31
51	GENOMIC ANALYSIS OF AVIAN INFLUENZA VIRUSES FROM WATERFOWL IN WESTERN ALASKA, USA. Journal of Wildlife Diseases, 2013, 49, 600-610.	0.8	26
52	Evidence that Life History Characteristics of Wild Birds Influence Infection and Exposure to Influenza A Viruses. PLoS ONE, 2013, 8, e57614.	2.5	26
53	Evaluation of Nobuto Filter Paper Strips for the Detection of Avian Influenza Virus Antibody in Waterfowl. Avian Diseases, 2011, 55, 674-676.	1.0	24
54	Developmental analysis and subcellular localization of the murine homologue of ELL. Proceedings of the National Academy of Sciences of the United States of America, 1997, 94, 1408-1413.	7.1	23

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55	High Rates of Detection of Clade 2.3.4.4 Highly Pathogenic Avian Influenza H5 Viruses in Wild Birds in the Pacific Northwest During the Winter of 2014–15. Avian Diseases, 2016, 60, 354-358.	1.0	21
56	Prevalence and Pathology of West Nile Virus in Naturally Infected House Sparrows, Western Nebraska, 2008. American Journal of Tropical Medicine and Hygiene, 2010, 82, 937-944.	1.4	20
57	PATHOLOGY AND VIRUS DETECTION IN TISSUES OF NESTLING HOUSE SPARROWS NATURALLY INFECTED WITH BUGGY CREEK VIRUS (TOGAVIRIDAE). Journal of Wildlife Diseases, 2010, 46, 23-32.	0.8	20
58	Evidence for limited exchange of avian influenza viruses between seaducks and dabbling ducks at Alaska Peninsula coastal lagoons. Archives of Virology, 2011, 156, 1813-1821.	2.1	20
59	Experimental challenge and pathology of highly pathogenic avian influenza virus H5N1 in dunlin (Calidris alpina), an intercontinental migrant shorebird species. Influenza and Other Respiratory Viruses, 2011, 5, 365-372.	3.4	19
60	Impacts of Migratory Sandhill Cranes (Grus canadensis) on Microbial Water Quality in the Central Platte River, Nebraska, USA. Water, Air, and Soil Pollution, 2013, 224, 1.	2.4	19
61	Expansion of an Exotic Species and Concomitant Disease Outbreaks: Pigeon Paramyxovirus in Free-Ranging Eurasian Collared Doves. EcoHealth, 2012, 9, 163-170.	2.0	18
62	Avian influenza virus ecology in Iceland shorebirds: Intercontinental reassortment and movement. Infection, Genetics and Evolution, 2014, 28, 130-136.	2.3	18
63	Surveillance for Highly Pathogenic Avian Influenza Virus in Wild Birds during Outbreaks in Domestic Poultry, Minnesota, 2015. Emerging Infectious Diseases, 2016, 22, 1278-1282.	4.3	18
64	Lethal Infection of Wild Raptors with Highly Pathogenic Avian Influenza H5N8 and H5N2 Viruses in the USA, 2014–15. Journal of Wildlife Diseases, 2019, 55, 164.	0.8	18
65	West Nile Virus Transmission in Winter: The 2013 Great Salt Lake Bald Eagle and Eared Grebes Mortality Event. PLOS Currents, 2014, 6, .	1.4	18
66	PATHOGEN EXPOSURE AND BLOOD CHEMISTRY IN THE WASHINGTON, USA POPULATION OF NORTHERN SEA OTTERS (<i>ENHYDRA LUTRIS KENYONI</i>). Journal of Wildlife Diseases, 2013, 49, 887-899.	0.8	17
67	Inferring epidemiologic dynamics from viral evolution: 2014–2015 Eurasian/North American highly pathogenic avian influenza viruses exceed transmission threshold, <i>R</i> ₀ Â=Â1, in wild birds and poultry in North America. Evolutionary Applications, 2018, 11, 547-557.	3.1	17
68	Serologic Evidence of Influenza A(H1N1)pdm09 Virus Infection in Northern Sea Otters. Emerging Infectious Diseases, 2014, 20, 915-917.	4.3	16
69	Demographic and Spatiotemporal Patterns of Avian Influenza Infection at the Continental Scale, and in Relation to Annual Life Cycle of a Migratory Host. PLoS ONE, 2015, 10, e0130662.	2.5	16
70	A picornavirus-like pathogen of Cotylogaster occidentalis (Trematoda: Aspidogastrea), an intestinal parasite of freshwater mollusks. Journal of Invertebrate Pathology, 1984, 43, 197-206.	3.2	15
71	The dynamics of avian influenza in Lesser Snow Geese: implications for annual and migratory infection patterns. Ecological Applications, 2015, 25, 1851-1859.	3.8	15
72	Detection of spring viraemia of carp virus in imported amphibians reveals an unanticipated foreign animal disease threat. Emerging Microbes and Infections, $2016, 5, 1-7$.	6.5	15

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73	A New Ranavirus Isolated fromPseudacris clarkiiTadpoles in Playa Wetlands in the Southern High Plains, Texas. Journal of Aquatic Animal Health, 2010, 22, 65-72.	1.4	14
74	The Effect of Swab Sample Choice on the Detection of Avian Influenza in Apparently Healthy Wild Ducks. Avian Diseases, 2012, 56, 114-119.	1.0	14
75	Identification of Two novel reassortant avian influenza a (H5N6) viruses in whooper swans in Korea, 2016. Virology Journal, 2017, 14, 60.	3.4	14
76	Artificial intelligence and avian influenza: Using machine learning to enhance active surveillance for avian influenza viruses. Transboundary and Emerging Diseases, 2019, 66, 2537-2545.	3.0	14
77	An mRNA-dependent in vitro translation system from Trypanosoma brucei. Molecular and Biochemical Parasitology, 1991, 46, 265-274.	1.1	13
78	Natural Infections With Pigeon Paramyxovirus Serotype 1: Pathologic Changes in Eurasian Collared-Doves (<i>Streptopelia decaocto</i>) and Rock Pigeons (<i>Columba livia</i>) in the United States. Veterinary Pathology, 2017, 54, 695-703.	1.7	11
79	Surveillance for Highly Pathogenic Avian Influenza in Wild Turkeys (Meleagris gallopavo) of Minnesota, USA during 2015 Outbreaks in Domestic Poultry. Journal of Wildlife Diseases, 2017, 53, 616-620.	0.8	11
80	A Blood Survey of Elements, Viral Antibodies, and Hemoparasites in Wintering Harlequin Ducks (Histrionicus Histrionicus) and Barrow's Goldeneyes (Bucephala Islandica). Journal of Wildlife Diseases, 2008, 44, 486-493.	0.8	10
81	Investigation of the 2018 thick-billed murre (Uria lomvia) die-off on St. Lawrence Island rules out food shortage as the cause. Deep-Sea Research Part II: Topical Studies in Oceanography, 2020, 181-182, 104879.	1.4	10
82	Experimental Challenge of a Peridomestic Avian Species, European Starlings (<i>Sturnus vulgaris</i>), with Novel Influenza A H7N9 Virus from China. Journal of Wildlife Diseases, 2016, 52, 709-712.	0.8	8
83	An Opportunistic Survey Reveals an Unexpected Coronavirus Diversity Hotspot in North America. Viruses, 2021, 13, 2016.	3 . 3	8
84	Avian influenza virus and free-ranging wild birds. Journal of the American Veterinary Medical Association, 2006, 228, 1877-1882.	0.5	7
85	Identification and characterization of Highlands J virus from a Mississippi sandhill crane using unbiased next-generation sequencing. Journal of Virological Methods, 2014, 206, 42-45.	2.1	7
86	SPATIAL AND TEMPORAL PATTERNS OF AVIAN PARAMYXOVIRUS-1 OUTBREAKS IN DOUBLE-CRESTED CORMORANTS (<i>PHALACROCORAX AURITUS</i>) IN THE USA. Journal of Wildlife Diseases, 2015, 51, 101-112.	0.8	7
87	Experimental Infection of Common Eider Ducklings with Wellfleet Bay Virus, a Newly Characterized Orthomyxovirus. Emerging Infectious Diseases, 2017, 23, 1958-1965.	4.3	7
88	Interlaboratory comparison of SARS-CoV2 molecular detection assays in use by U.S. veterinary diagnostic laboratories. Journal of Veterinary Diagnostic Investigation, 2021, 33, 1039-1051.	1.1	7
89	Total Protein Concentration and Diagnostic Test Results for Gray Wolf (Canis lupus) Serum using Nobuto Filter Paper Strips. Journal of Wildlife Diseases, 2015, 51, 475-478.	0.8	4
90	Emergence and molecular characterization of pigeon Paramyxovirus-1 in non-native Eurasian collared doves (Streptopelia decaocto) in California, USA. Infection, Genetics and Evolution, 2021, 91, 104809.	2.3	4

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91	Validation of a Real-Time Reverse Transcriptase–PCR Assay for the Detection of H7 Avian Influenza Virus. Avian Diseases, 2010, 54, 639-643.	1.0	3
92	H9N2 Influenza A Virus Isolated from a Greater White-Fronted Wild Goose (Anser albifrons) in Alaska Has a Mutation in the PB2 Gene, Which Is Associated with Pathogenicity in Human Pandemic 2009 H1N1. Genome Announcements, 2016 , 4 , $.$	0.8	3
93	Changes in West Nile Virus Seroprevalence and Antibody Titers among Wisconsin Mesopredators 2003–2006. American Journal of Tropical Medicine and Hygiene, 2009, 81, 177-179.	1.4	2
94	Inactivation of Viable Surrogates for the Select Agents Virulent Newcastle Disease Virus and Highly Pathogenic Avian Influenza Virus Using Either Commercial Lysis Buffer or Heat. Applied Biosafety, 2019, 24, 189-199.	0.5	1