Daniel R. Perez

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Intra- and inter-host evolution of H9N2 influenza A virus in Japanese quail. Virus Evolution, 2022, 8, veac001.	2.2	8
2	Reverse Genetics for Influenza A and B Viruses Driven by Swine Polymerase I Promoter. Methods in Molecular Biology, 2022, 2465, 257-281.	0.4	1
3	Infectious Salmon Anemia Virus Infectivity Is Determined by Multiple Segments with an Important Contribution from Segment 5. Viruses, 2022, 14, 631.	1.5	5
4	Influenza antivirals and animal models. FEBS Open Bio, 2022, 12, 1142-1165.	1.0	18
5	Naturally Acquired Antibodies to Influenza A Virus in Fall-Migrating North American Mallards. Veterinary Sciences, 2022, 9, 214.	0.6	2
6	Robustness of the Ferret Model for Influenza Risk Assessment Studies: a Cross-Laboratory Exercise. MBio, 2022, 13, .	1.8	12
7	Development of a swine RNA polymerase I driven Influenza reverse genetics system for the rescue of type A and B Influenza viruses. Journal of Virological Methods, 2021, 288, 114011.	1.0	3
8	Mutations in PB1, NP, HA, and NA Contribute to Increased Virus Fitness of H5N2 Highly Pathogenic Avian Influenza Virus Clade 2.3.4.4 in Chickens. Journal of Virology, 2021, 95, .	1.5	11
9	Rational design of a deuterium-containing M2-S31N channel blocker UAWJ280 with <i>in vivo</i> antiviral efficacy against both oseltamivir sensitive and -resistant influenza A viruses. Emerging Microbes and Infections, 2021, 10, 1832-1848.	3.0	10
10	Efficacy of GC-376 against SARS-CoV-2 virus infection in the K18 hACE2 transgenic mouse model. Scientific Reports, 2021, 11, 9609.	1.6	46
11	Development of a Novel Live Attenuated Influenza A Virus Vaccine Encoding the IgA-Inducing Protein. Vaccines, 2021, 9, 703.	2.1	8
12	A New Master Donor Virus for the Development of Live-Attenuated Influenza B Virus Vaccines. Viruses, 2021, 13, 1278.	1.5	2
13	Mutation E48K in PB1 Polymerase Subunit Improves Stability of a Candidate Live Attenuated Influenza B Virus Vaccine. Vaccines, 2021, 9, 800.	2.1	4
14	FluB-RAM and FluB-RANS: Genome Rearrangement as Safe and Efficacious Live Attenuated Influenza B Virus Vaccines. Vaccines, 2021, 9, 897.	2.1	2
15	2021 Taxonomic update of phylum Negarnaviricota (Riboviria: Orthornavirae), including the large orders Bunyavirales and Mononegavirales. Archives of Virology, 2021, 166, 3513-3566.	0.9	62
16	Airborne Transmission of Avian Origin H9N2 Influenza A Viruses in Mammals. Viruses, 2021, 13, 1919.	1.5	19
17	Evolution and Antigenic Advancement of N2 Neuraminidase of Swine Influenza A Viruses Circulating in the United States following Two Separate Introductions from Human Seasonal Viruses. Journal of Virology, 2021, 95, e0063221.	1.5	10
18	Mild and Severe SARS-CoV-2 Infection Induces Respiratory and Intestinal Microbiome Changes in the K18-hACE2 Transgenic Mouse Model. Microbiology Spectrum, 2021, 9, e0053621.	1.2	21

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19	H9 Influenza Viruses: An Emerging Challenge. Cold Spring Harbor Perspectives in Medicine, 2020, 10, a038588.	2.9	58
20	Reverse genetics for influenza B viruses and recent advances in vaccine development. Current Opinion in Virology, 2020, 44, 191-202.	2.6	7
21	2020 taxonomic update for phylum Negarnaviricota (Riboviria: Orthornavirae), including the large orders Bunyavirales and Mononegavirales. Archives of Virology, 2020, 165, 3023-3072.	0.9	184
22	The pneumococcal two-component system SirRH is linked to enhanced intracellular survival of Streptococcus pneumoniae in influenza-infected pulmonary cells. PLoS Pathogens, 2020, 16, e1008761.	2.1	11
23	Silent Infection of B and CD8 + T Lymphocytes by Influenza A Virus in Children with Tonsillar Hypertrophy. Journal of Virology, 2020, 94, .	1.5	5
24	Editorial: Emerging Swine Viruses. Frontiers in Veterinary Science, 2020, 7, 132.	0.9	8
25	Collective interactions augment influenza A virus replication in a host-dependent manner. Nature Microbiology, 2020, 5, 1158-1169.	5.9	32
26	In vivo rescue of recombinant Zika virus from an infectious cDNA clone and its implications in vaccine development. Scientific Reports, 2020, 10, 512.	1.6	14
27	Characterizing Emerging Canine H3 Influenza Viruses. PLoS Pathogens, 2020, 16, e1008409.	2.1	29
28	Plasmid-Based Reverse Genetics of Influenza A Virus. Methods in Molecular Biology, 2020, 2123, 37-59.	0.4	10
29	LIMITED DETECTION OF ANTIBODIES TO CLADE 2.3.4.4 A/GOOSE/GUANGDONG/1/1996 LINEAGE HIGHLY PATHOGENIC H5 AVIAN INFLUENZA VIRUS IN NORTH AMERICAN WATERFOWL. Journal of Wildlife Diseases, 2020, 56, 47.	0.3	6
30	LIMITED DETECTION OF ANTIBODIES TO CLADE 2.3.4.4 A/GOOSE/GUANGDONG/1/1996 LINEAGE HIGHLY PATHOGENIC H5 AVIAN INFLUENZA VIRUS IN NORTH AMERICAN WATERFOWL. Journal of Wildlife Diseases, 2020, 56, 47-57.	0.3	1
31	Characterizing Emerging Canine H3 Influenza Viruses. , 2020, 16, e1008409.		0
32	Characterizing Emerging Canine H3 Influenza Viruses. , 2020, 16, e1008409.		0
33	Characterizing Emerging Canine H3 Influenza Viruses. , 2020, 16, e1008409.		0
34	Characterizing Emerging Canine H3 Influenza Viruses. , 2020, 16, e1008409.		0
35	Characterizing Emerging Canine H3 Influenza Viruses. , 2020, 16, e1008409.		0
36	Characterizing Emerging Canine H3 Influenza Viruses. , 2020, 16, e1008409.		0

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37	Improved detection of influenza A virus from blueâ€winged teals by sequencing directly from swab material. Ecology and Evolution, 2019, 9, 6534-6546.	0.8	18
38	Maternally-Derived Antibodies Protect against Challenge with Highly Pathogenic Avian Influenza Virus of the H7N3 Subtype. Vaccines, 2019, 7, 163.	2.1	4
39	Age-dependent pathogenesis of clade 2.3.4.4A H5N2 HPAIV in experimentally infected Broad Breasted White turkeys. Veterinary Microbiology, 2019, 231, 183-190.	0.8	5
40	Flexibility <i>In Vitro</i> of Amino Acid 226 in the Receptor-Binding Site of an H9 Subtype Influenza A Virus and Its Effect <i>In Vivo</i> on Virus Replication, Tropism, and Transmission. Journal of Virology, 2019, 93, .	1.5	34
41	Plasticity of Amino Acid Residue 145 Near the Receptor Binding Site of H3 Swine Influenza A Viruses and Its Impact on Receptor Binding and Antibody Recognition. Journal of Virology, 2019, 93, .	1.5	19
42	Enhancing the cross protective efficacy of live attenuated influenza virus vaccine by supplemented vaccination with M2 ectodomain virus-like particles. Virology, 2019, 529, 111-121.	1.1	15
43	Avian Influenza Virus. , 2019, , .		1
44	Evidence of a fixed internal gene constellation in influenza A viruses isolated from wild birds in Argentina (2006–2016). Emerging Microbes and Infections, 2018, 7, 1-13.	3.0	15
45	Identification of Amino Acid Residues Responsible for Inhibition of Host Gene Expression by Influenza A H9N2 NS1 Targeting of CPSF30. Frontiers in Microbiology, 2018, 9, 2546.	1.5	15
46	Comparison of Adjuvanted-Whole Inactivated Virus and Live-Attenuated Virus Vaccines against Challenge with Contemporary, Antigenically Distinct H3N2 Influenza A Viruses. Journal of Virology, 2018, 92, .	1.5	11
47	Universal Vaccines and Vaccine Platforms to Protect against Influenza Viruses in Humans and Agriculture. Frontiers in Microbiology, 2018, 9, 123.	1.5	108
48	Alternative Strategy for a Quadrivalent Live Attenuated Influenza Virus Vaccine. Journal of Virology, 2018, 92, .	1.5	10
49	Crosstalk between the serine/threonine kinase StkP and the response regulator ComE controls the stress response and intracellular survival of Streptococcus pneumoniae. PLoS Pathogens, 2018, 14, e1007118.	2.1	33
50	Adaptation of Human Influenza Viruses to Swine. Frontiers in Veterinary Science, 2018, 5, 347.	0.9	61
51	Two years of surveillance of influenza a virus infection in a swine herd. Results of virological, serological and pathological studies. Comparative Immunology, Microbiology and Infectious Diseases, 2017, 50, 110-115.	0.7	7
52	Widespread Virus Replication in Alveoli Drives Acute Respiratory Distress Syndrome in Aerosolized H5N1 Influenza Infection of Macaques. Journal of Immunology, 2017, 198, 1616-1626.	0.4	40
53	Factors affecting induction of peripheral IFN-Î ³ recall response to influenza A virus vaccination in pigs. Veterinary Immunology and Immunopathology, 2017, 185, 57-65.	0.5	15
54	Plasmid-Based Reverse Genetics of Influenza A Virus. Methods in Molecular Biology, 2017, 1602, 251-273.	0.4	22

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55	Influenza A virus vaccines for swine. Veterinary Microbiology, 2017, 206, 35-44.	0.8	85
56	Reverse Genetics of Influenza B Viruses. Methods in Molecular Biology, 2017, 1602, 205-238.	0.4	21
57	Development of an Alternative Modified Live Influenza B Virus Vaccine. Journal of Virology, 2017, 91, .	1.5	17
58	Distribution of O-Acetylated Sialic Acids among Target Host Tissues for Influenza Virus. MSphere, 2017, 2, .	1.3	56
59	Short- and long-term protective efficacy against clade 2.3.4.4 H5N2 highly pathogenic avian influenza virus following prime-boost vaccination in turkeys. Vaccine, 2017, 35, 5637-5643.	1.7	22
60	Replication of H9 influenza viruses in the human ex vivo respiratory tract, and the influence of neuraminidase on virus release. Scientific Reports, 2017, 7, 6208.	1.6	7
61	Evidence of Intercontinental Spread and Uncommon Variants of Low-Pathogenicity Avian Influenza Viruses in Ducks Overwintering in Guatemala. MSphere, 2017, 2, .	1.3	8
62	Origin, distribution, and potential risk factors associated with influenza A virus in swine in two production systems in Guatemala. Influenza and Other Respiratory Viruses, 2017, 11, 182-192.	1.5	13
63	Age at Vaccination and Timing of Infection Do Not Alter Vaccine-Associated Enhanced Respiratory Disease in Influenza A Virus-Infected Pigs. Vaccine Journal, 2016, 23, 470-482.	3.2	19
64	Impact of a potential glycosylation site at neuraminidase amino acid 264 of influenza A/H9N2 virus. Veterinary Microbiology, 2016, 196, 9-13.	0.8	3
65	The Molecular Determinants of Antibody Recognition and Antigenic Drift in the H3 Hemagglutinin of Swine Influenza A Virus. Journal of Virology, 2016, 90, 8266-8280.	1.5	54
66	Prevalence and Diversity of Low Pathogenicity Avian Influenza Viruses in Wild Birds in Guatemala, 2010–2013. Avian Diseases, 2016, 60, 359-364.	0.4	17
67	Neuraminidase inhibiting antibody responses in pigs differ between influenza A virus N2 lineages and by vaccine type. Vaccine, 2016, 34, 3773-3779.	1.7	12
68	Replication-competent fluorescent-expressing influenza B virus. Virus Research, 2016, 213, 69-81.	1.1	37
69	Heterologous challenge in the presence of maternally-derived antibodies results in vaccine-associated enhanced respiratory disease in weaned piglets. Virology, 2016, 491, 79-88.	1.1	25
70	Vaccine-associated enhanced respiratory disease is influenced by haemagglutinin and neuraminidase in whole inactivated influenza virus vaccines. Journal of General Virology, 2016, 97, 1489-1499.	1.3	46
71	Replication-Competent Influenza A and B Viruses Expressing a Fluorescent Dynamic Timer Protein for In Vitro and In Vivo Studies. PLoS ONE, 2016, 11, e0147723.	1.1	32
72	Replication and transmission of mammalian-adapted H9 subtype influenza virus in pigs and quail. Journal of General Virology, 2015, 96, 2511-2521.	1.3	14

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73	An efficient and rapid influenza gene cloning strategy for reverse genetics system. Journal of Virological Methods, 2015, 222, 91-94.	1.0	14
74	Oral Fluids as a Live-Animal Sample Source for Evaluating Cross-Reactivity and Cross-Protection following Intranasal Influenza A Virus Vaccination in Pigs. Vaccine Journal, 2015, 22, 1109-1120.	3.2	14
75	Novel Reassortant Human-Like H3N2 and H3N1 Influenza A Viruses Detected in Pigs Are Virulent and Antigenically Distinct from Swine Viruses Endemic to the United States. Journal of Virology, 2015, 89, 11213-11222.	1.5	84
76	Genomic Characterization of H14 Subtype Influenza A Viruses in New World Waterfowl and Experimental Infectivity in Mallards (Anas platyrhynchos). PLoS ONE, 2014, 9, e95620.	1.1	23
77	Modeling Human Respiratory Viral Infections in the Cotton Rat (Sigmodon hispidus). Journal of Antivirals & Antiretrovirals, 2014, 06, 40-42.	0.1	20
78	Antigenic Variation of Clade 2.1 H5N1 Virus Is Determined by a Few Amino Acid Substitutions Immediately Adjacent to the Receptor Binding Site. MBio, 2014, 5, e01070-14.	1.8	57
79	Evidence for Seasonal Patterns in the Relative Abundance of Avian Influenza Virus Subtypes in Blue-Winged Teal (Anas discors). Journal of Wildlife Diseases, 2014, 50, 916-922.	0.3	36
80	Polymorphisms in the haemagglutinin gene influenced the viral shedding of pandemic 2009 influenza virus in swine. Journal of General Virology, 2014, 95, 2618-2626.	1.3	4
81	Genome rearrangement of influenza virus for anti-viral drug screening. Virus Research, 2014, 189, 14-23.	1.1	22
82	Airborne Transmission of Highly Pathogenic H7N1 Influenza Virus in Ferrets. Journal of Virology, 2014, 88, 6623-6635.	1.5	83
83	Alternative Reassortment Events Leading to Transmissible H9N1 Influenza Viruses in the Ferret Model. Journal of Virology, 2014, 88, 66-71.	1.5	36
84	Live attenuated influenza A virus vaccine protects against A(H1N1)pdm09 heterologous challenge without vaccine associated enhanced respiratory disease. Virology, 2014, 471-473, 93-104.	1.1	60
85	Correction for Koel et al., Antigenic Variation of Clade 2.1 H5N1 Virus Is Determined by a Few Amino Acid Substitutions Immediately Adjacent to the Receptor Binding Site. MBio, 2014, 5, .	1.8	6
86	Influenza A and B Virus Intertypic Reassortment through Compatible Viral Packaging Signals. Journal of Virology, 2014, 88, 10778-10791.	1.5	83
87	All-in-One Bacmids: an Efficient Reverse Genetics Strategy for Influenza A Virus Vaccines. Journal of Virology, 2014, 88, 10013-10025.	1.5	20
88	Design of Alternative Live Attenuated Influenza Virus Vaccines. Current Topics in Microbiology and Immunology, 2014, 386, 205-235.	0.7	6
89	Infection and Pathogenesis of Canine, Equine, and Human Influenza Viruses in Canine Tracheas. Journal of Virology, 2014, 88, 9208-9219.	1.5	37
90	Interactions between the Influenza A Virus RNA Polymerase Components and Retinoic Acid-Inducible Gene I. Journal of Virology, 2014, 88, 10432-10447.	1.5	38

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91	Roles of Major Histocompatibility Complex Class II in Inducing Protective Immune Responses to Influenza Vaccination. Journal of Virology, 2014, 88, 7764-7775.	1.5	23
92	Gain-of-function experiments on H7N9. Nature, 2013, 500, 150-151.	13.7	24
93	MicroRNA-based strategy to mitigate the risk of gain-of-function influenza studies. Nature Biotechnology, 2013, 31, 844-847.	9.4	77
94	Receptor Characterization and Susceptibility of Cotton Rats to Avian and 2009 Pandemic Influenza Virus Strains. Journal of Virology, 2013, 87, 2036-2045.	1.5	34
95	H5N1, a wealth of knowledge to improve pandemic preparedness. Virus Research, 2013, 178, 1-2.	1.1	10
96	The effect of avian influenza virus NS1 allele on virus replication and innate gene expression in avian cells. Molecular Immunology, 2013, 56, 358-368.	1.0	25
97	Evidence for avian H9N2 influenza virus infections among rural villagers in Cambodia. Journal of Infection and Public Health, 2013, 6, 69-79.	1.9	46
98	Influenza Viruses with Rearranged Genomes as Live-Attenuated Vaccines. Journal of Virology, 2013, 87, 5118-5127.	1.5	57
99	Transmission Studies Resume for Avian Flu. Science, 2013, 339, 520-521.	6.0	34
100	Amino Acid 316 of Hemagglutinin and the Neuraminidase Stalk Length Influence Virulence of H9N2 Influenza Virus in Chickens and Mice. Journal of Virology, 2013, 87, 2963-2968.	1.5	70
101	Molecular Basis for Broad Neuraminidase Immunity: Conserved Epitopes in Seasonal and Pandemic H1N1 as Well as H5N1 Influenza Viruses. Journal of Virology, 2013, 87, 9290-9300.	1.5	141
102	Efficacy in Pigs of Inactivated and Live Attenuated Influenza Virus Vaccines against Infection and Transmission of an Emerging H3N2 Similar to the 2011-2012 H3N2v. Journal of Virology, 2013, 87, 9895-9903.	1.5	88
103	Gain-of-Function Experiments on H7N9. Science, 2013, 341, 612-613.	6.0	24
104	In Vivo Selection of H1N2 Influenza Virus Reassortants in the Ferret Model. Journal of Virology, 2013, 87, 3277-3283.	1.5	12
105	Swine influenza virus vaccine serologic crossâ€reactivity to contemporary <scp>US</scp> swine H3N2 and efficacy in pigs infected with an H3N2 similar to 2011–2012 H3N2v. Influenza and Other Respiratory Viruses, 2013, 7, 32-41.	1.5	34
106	Swine influenza: clinical, serological, pathological, and virological crossâ€sectional studies in nine farms in <scp>A</scp> rgentina. Influenza and Other Respiratory Viruses, 2013, 7, 10-15.	1.5	12
107	Influenza A(H1N1)pdm09 virus infection in marine mammals in California. Emerging Microbes and Infections, 2013, 2, 1-2.	3.0	11
108	The non-structural (NS) gene segment of H9N2 influenza virus isolated from backyard poultry in Pakistan reveals strong genetic and functional similarities to the NS gene of highly pathogenic H5N1. Virulence, 2013, 4, 612-623.	1.8	12

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109	Deletions in the Neuraminidase Stalk Region of H2N2 and H9N2 Avian Influenza Virus Subtypes Do Not Affect Postinfluenza Secondary Bacterial Pneumonia. Journal of Virology, 2012, 86, 3564-3573.	1.5	19
110	Pause on Avian Flu Transmission Research. Science, 2012, 335, 400-401.	6.0	58
111	H5N1 Debates: Hung Up on the Wrong Questions. Science, 2012, 335, 799-801.	6.0	14
112	Restored PB1-F2 in the 2009 Pandemic H1N1 Influenza Virus Has Minimal Effects in Swine. Journal of Virology, 2012, 86, 5523-5532.	1.5	33
113	Isolation and characterization of an H9N2 influenza virus isolated in Argentina. Virus Research, 2012, 168, 41-47.	1.1	27
114	Where Do Avian Influenza Viruses Meet in the Americas?. Avian Diseases, 2012, 56, 1025-1033.	0.4	22
115	Heightened adaptive immune responses following vaccination with a temperature-sensitive, live-attenuated influenza virus compared to adjuvanted, whole-inactivated virus in pigs. Vaccine, 2012, 30, 5830-5838.	1.7	40
116	Strain-dependent effects of PB1-F2 of triple-reassortant H3N2 influenza viruses in swine. Journal of General Virology, 2012, 93, 2204-2214.	1.3	21
117	Influenza A Viruses from Wild Birds in Guatemala Belong to the North American Lineage. PLoS ONE, 2012, 7, e32873.	1.1	39
118	Partial and Full PCR-Based Reverse Genetics Strategy for Influenza Viruses. PLoS ONE, 2012, 7, e46378.	1.1	22
119	Passive immune neutralization strategies for prevention and control of influenza A infections. Immunotherapy, 2012, 4, 175-186.	1.0	17
120	Cold-Adapted Influenza and Recombinant Adenovirus Vaccines Induce Cross-Protective Immunity against pH1N1 Challenge in Mice. PLoS ONE, 2011, 6, e21937.	1.1	42
121	Modifications in the Polymerase Genes of a Swine-Like Triple-Reassortant Influenza Virus To Generate Live Attenuated Vaccines against 2009 Pandemic H1N1 Viruses. Journal of Virology, 2011, 85, 456-469.	1.5	85
122	Evidence of reassortment of pandemic H1N1 influenza virus in swine in Argentina: are we facing the expansion of potential epicenters of influenza emergence?. Influenza and Other Respiratory Viruses, 2011, 5, 409-412.	1.5	49
123	A novel monoclonal antibody effective against lethal challenge with swine-lineage and 2009 pandemic H1N1 influenza viruses in mice. Virology, 2011, 417, 379-384.	1.1	8
124	Improved hatchability and efficient protection after in ovo vaccination with live-attenuated H7N2 and H9N2 avian influenza viruses. Virology Journal, 2011, 8, 31.	1.4	19
125	GM-CSF in the Lung Protects against Lethal Influenza Infection. American Journal of Respiratory and Critical Care Medicine, 2011, 184, 259-268.	2.5	139
126	Outbreak of swine influenza in Argentina reveals a non-contemporary human H3N2 virus highly transmissible among pigs. Journal of General Virology, 2011, 92, 2871-2878.	1.3	39

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127	Influenza Virus Aerosols in Human Exhaled Breath: Particle Size, Culturability, and Effect of Surgical Masks. Epidemiology, 2011, 22, S51.	1.2	8
128	Seasonal FluMist Vaccination Induces Cross-Reactive T Cell Immunity against H1N1 (2009) Influenza and Secondary Bacterial Infections. Journal of Immunology, 2011, 186, 987-993.	0.4	83
129	Phylogenetic Analysis of H6 Influenza Viruses Isolated from Rosy-Billed Pochards (Netta peposaca) in Argentina Reveals the Presence of Different HA Gene Clusters. Journal of Virology, 2011, 85, 13354-13362.	1.5	27
130	Compatibility of H9N2 avian influenza surface genes and 2009 pandemic H1N1 internal genes for transmission in the ferret model. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 12084-12088.	3.3	117
131	Differential Contribution of PB1-F2 to the Virulence of Highly Pathogenic H5N1 Influenza A Virus in Mammalian and Avian Species. PLoS Pathogens, 2011, 7, e1002186.	2.1	119
132	A monoclonal antibody-based ELISA for differential diagnosis of 2009 pandemic H1N1. Influenza and Other Respiratory Viruses, 2011, 5, 138-41.	1.5	0
133	Influenza Virus Aerosols In Human Exhaled Breath: Particle Size, Culturability, And Effect Of Surgical Masks. , 2010, , .		5
134	Introduction of Virulence Markers in PB2 of Pandemic Swine-Origin Influenza Virus Does Not Result in Enhanced Virulence or Transmission. Journal of Virology, 2010, 84, 3752-3758.	1.5	126
135	Contributions of the Avian Influenza Virus HA, NA, and M2 Surface Proteins to the Induction of Neutralizing Antibodies and Protective Immunity. Journal of Virology, 2010, 84, 2408-2420.	1.5	59
136	Virulence-Associated Substitution D222G in the Hemagglutinin of 2009 Pandemic Influenza A(H1N1) Virus Affects Receptor Binding. Journal of Virology, 2010, 84, 11802-11813.	1.5	197
137	Mutations in the NS1 C-terminal tail do not enhance replication or virulence of the 2009 pandemic H1N1 influenza A virus. Journal of General Virology, 2010, 91, 1737-1742.	1.3	58
138	Variations in the Hemagglutinin of the 2009 H1N1 Pandemic Virus: Potential for Strains with Altered Virulence Phenotype?. PLoS Pathogens, 2010, 6, e1001145.	2.1	103
139	Inefficient Control of Host Gene Expression by the 2009 Pandemic H1N1 Influenza A Virus NS1 Protein. Journal of Virology, 2010, 84, 6909-6922.	1.5	152
140	Alternative Live-Attenuated Influenza Vaccines Based on Modifications in the Polymerase Genes Protect against Epidemic and Pandemic Flu. Journal of Virology, 2010, 84, 4587-4596.	1.5	41
141	Intranasal Delivery of an IgA Monoclonal Antibody Effective against Sublethal H5N1 Influenza Virus Infection in Mice. Vaccine Journal, 2010, 17, 1363-1370.	3.2	36
142	A 27-Amino-Acid Deletion in the Neuraminidase Stalk Supports Replication of an Avian H2N2 Influenza A Virus in the Respiratory Tract of Chickens. Journal of Virology, 2010, 84, 11831-11840.	1.5	69
143	Characterization of influenza virus sialic acid receptors in minor poultry species. Virology Journal, 2010, 7, 365.	1.4	81
144	Adaptation of a Mallard H5N2 Low Pathogenicity Influenza Virus in Chickens with Prior History of Infection with Infectious Bursal Disease Virus. Avian Diseases, 2010, 54, 513-521.	0.4	20

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145	Immunization of Chickens with Newcastle Disease Virus Expressing H5 Hemagglutinin Protects against Highly Pathogenic H5N1 Avian Influenza Viruses. PLoS ONE, 2009, 4, e6509.	1.1	70
146	Differential regulation of antiviral and proinflammatory cytokines and suppression of Fas-mediated apoptosis by NS1 of H9N2 avian influenza virus in chicken macrophages. Journal of General Virology, 2009, 90, 1109-1118.	1.3	36
147	Live Attenuated Influenza Viruses Containing NS1 Truncations as Vaccine Candidates against H5N1 Highly Pathogenic Avian Influenza. Journal of Virology, 2009, 83, 1742-1753.	1.5	217
148	The prevention and control of avian influenza: The avian influenza coordinated agriculture project. Poultry Science, 2009, 88, 837-841.	1.5	2
149	Partial direct contact transmission in ferrets of a mallard H7N3 influenza virus with typical avian-like receptor specificity. Virology Journal, 2009, 6, 126.	1.4	22
150	Minimal molecular constraints for respiratory droplet transmission of an avian–human H9N2 influenza A virus. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 7565-7570.	3.3	202
151	Fitness of Pandemic H1N1 and Seasonal influenza A viruses during Co-infectionEvidence of competitive advantage of pandemic H1N1 influenza versus seasonal influenza. PLOS Currents, 2009, 1, RRN1011.	1.4	30
152	Avian influenza virus isolated in wild waterfowl in Argentina: Evidence of a potentially unique phylogenetic lineage in South America. Virology, 2008, 378, 363-370.	1.1	82
153	An avian live attenuated master backbone for potential use in epidemic and pandemic influenza vaccines. Journal of General Virology, 2008, 89, 2682-2690.	1.3	23
154	A new influenza virus virulence determinant: The NS1 protein four C-terminal residues modulate pathogenicity. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 4381-4386.	3.3	375
155	Multivalent HA DNA Vaccination Protects against Highly Pathogenic H5N1 Avian Influenza Infection in Chickens and Mice. PLoS ONE, 2008, 3, e2432.	1.1	46
156	Replication and Transmission of H9N2 Influenza Viruses in Ferrets: Evaluation of Pandemic Potential. PLoS ONE, 2008, 3, e2923.	1.1	248
157	Evidence of Expanded Host Range and Mammalian-Associated Genetic Changes in a Duck H9N2 Influenza Virus Following Adaptation in Quail and Chickens. PLoS ONE, 2008, 3, e3170.	1.1	116
158	Pandemic Influenza: Preventing the Emergence of Novel Strains and Countermeasures to Ameliorate its Effects. Infectious Disorders - Drug Targets, 2007, 7, 304-317.	0.4	3
159	Genesis of pandemic influenza. Cytogenetic and Genome Research, 2007, 117, 394-402.	0.6	12
160	A New Generation of Modified Live-Attenuated Avian Influenza Viruses Using a Two-Strategy Combination as Potential Vaccine Candidates. Journal of Virology, 2007, 81, 9238-9248.	1.5	84
161	Immunosenescence and age-related susceptibility to influenza virus in Japanese quail. Developmental and Comparative Immunology, 2007, 31, 407-414.	1.0	30
162	Adaptation of Influenza A/Mallard/Potsdam/178-4/83 H2N2 Virus in Japanese Quail Leads to Infection and Transmission in Chickens. Avian Diseases, 2007, 51, 264-268.	0.4	47

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