

Xiufan Liu

List of Publications by Year in descending order

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229
papers

4,605
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136950

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232
docs citations

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4821
citing authors

#	ARTICLE	IF	CITATIONS
1	Extended state observer-based nonsingular terminal sliding mode controller for a DC-DC buck converter with disturbances: theoretical analysis and experimental verification. <i>International Journal of Control</i> , 2023, 96, 1661-1671.	1.9	8
2	Phylogenetic analysis and pathogenicity assessment of pigeon paramyxovirus type 1 circulating in China during 2007–2019. <i>Transboundary and Emerging Diseases</i> , 2022, 69, 2076-2088.	3.0	4
3	Emergence of a novel reassortant avian influenza virus (H10N3) in Eastern China with high pathogenicity and respiratory droplet transmissibility to mammals. <i>Science China Life Sciences</i> , 2022, 65, 1024-1035.	4.9	20
4	ESO-Based Vibration Control for All-Clamped Plate Using an Electrodynamics Inertial Actuator. <i>International Journal of Structural Stability and Dynamics</i> , 2022, 22, .	2.4	5
5	Newcastle disease virus degrades SIRT3 via PINK1-PRKN-dependent mitophagy to reprogram energy metabolism in infected cells. <i>Autophagy</i> , 2022, 18, 1503-1521.	9.1	52
6	Development of an indirect ELISA method based on the VP4 protein for detection antibody against duck hepatitis A virus type 1. <i>Journal of Virological Methods</i> , 2022, 300, 114393.	2.1	1
7	Baculovirus-derived influenza virus-like particle confers complete protection against lethal H7N9 avian influenza virus challenge in chickens and mice. <i>Veterinary Microbiology</i> , 2022, 264, 109306.	1.9	4
8	Novel reassortment 2.3.4.4b H5N8 highly pathogenic avian influenza viruses circulating in Xinjiang, China. <i>Preventive Veterinary Medicine</i> , 2022, 199, 105564.	1.9	2
9	Phylogenetic and phenotypic characterization of two novel clade 2.3.2.1 H5N2 subtype avian influenza viruses from chickens in China. <i>Infection, Genetics and Evolution</i> , 2022, 98, 105205.	2.3	3
10	Expression and characterization of a recombinant broadly-reactive monoclonal antibody against group 1 and 2 influenza viruses. <i>Protein Expression and Purification</i> , 2022, 192, 106046.	1.3	0
11	Effects of HA2 154 deglycosylation and NA V202I mutation on biological property of H5N6 subtype avian influenza virus. <i>Veterinary Microbiology</i> , 2022, 266, 109353.	1.9	2
12	Reduced-Order Extended State Observer-Based Sliding Mode Control for All-Clamped Plate Using an Inertial Actuator. <i>Energies</i> , 2022, 15, 1780.	3.1	4
13	Characterization of antibody response to an epitope spanning the haemagglutinin cleavage site of H7N9 subtype avian influenza virus for differentiation of infected and vaccinated chickens. <i>Avian Pathology</i> , 2022, , 1-25.	2.0	0
14	Nonlinear ESO-based vibration control for an all-clamped piezoelectric plate with disturbances and time delay: Design and hardware implementation. <i>Journal of Intelligent Material Systems and Structures</i> , 2022, 33, 2321-2335.	2.5	7
15	Generation of an avian influenza DIVA vaccine with a H3-peptide replacement located at HA2 against both highly and low pathogenic H7N9 virus. <i>Virulence</i> , 2022, 13, 530-541.	4.4	2
16	H5N1 infection impairs the alveolar epithelial barrier through intercellular junction proteins via Itch-mediated proteasomal degradation. <i>Communications Biology</i> , 2022, 5, 186.	4.4	9
17	Characterization of two chicken origin highly pathogenic H7N9 viruses isolated in northern China. <i>Veterinary Microbiology</i> , 2022, 268, 109394.	1.9	4
18	Emerging of H5N6 Subtype Influenza Virus with 129-Glycosylation Site on Hemagglutinin in Poultry in China Acquires Immune Pressure Adaption. <i>Microbiology Spectrum</i> , 2022, 10, e0253721.	3.0	3

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19	Single and Combined Effects of <i>Clostridium butyricum</i> and Coccidiosis Vaccine on Growth Performance and the Intestinal Microbiome of Broiler Chickens. <i>Frontiers in Microbiology</i> , 2022, 13, 811428.	3.5	5
20	Intranasal Immunization with a Recombinant Avian Paramyxovirus Serotypes 2 Vector-Based Vaccine Induces Protection against H9N2 Avian Influenza in Chicken. <i>Viruses</i> , 2022, 14, 918.	3.3	3
21	Experimental Study of Structural Vibration Control Based on Piezoelectric Shunt Control System. , 2022, , .		0
22	Genome-Wide Reassortment Analysis of Influenza A H7N9 Viruses Circulating in China during 2013â€“2019. <i>Viruses</i> , 2022, 14, 1256.	3.3	2
23	H5N1 avian influenza virus without 80â€“84 amino acid deletion at the NS1 protein hijacks the innate immune system of dendritic cells for an enhanced mammalian pathogenicity. <i>Transboundary and Emerging Diseases</i> , 2021, 68, 2401-2413.	3.0	9
24	Fabrication of chondroitin sulfate calcium complex and its chondrocyte proliferation in vitro. <i>Carbohydrate Polymers</i> , 2021, 254, 117282.	10.2	20
25	Phylogenetic tracing and biological characterization of a novel clade 2.3.2.1 reassortant of H5N6 subtype avian influenza virus in China. <i>Transboundary and Emerging Diseases</i> , 2021, 68, 730-741.	3.0	6
26	Pathogenicity and transmissibility of an H9N2 avian influenza virus that naturally harbors the mammalian-adaptive molecular factors in the hemagglutinin and PB2 proteins. <i>Journal of Infection</i> , 2021, 82, e22-e23.	3.3	10
27	Deep sequencing of the transcriptome from murine lung infected with H5N8 subtype avian influenza virus with combined substitutions I283M and K526R in PB2 gene. <i>Infection, Genetics and Evolution</i> , 2021, 87, 104672.	2.3	3
28	Novel reassortant 2.3.4.4B H5N6 highly pathogenic avian influenza viruses circulating among wild, domestic birds in Xinjiang, Northwest China. <i>Journal of Veterinary Science</i> , 2021, 22, e43.	1.3	4
29	Re-emergence of H5N8 highly pathogenic avian influenza virus in wild birds, China. <i>Emerging Microbes and Infections</i> , 2021, 10, 1819-1823.	6.5	17
30	Identification and Characterization of the ATG8, a Marker of <i>Eimeria tenella</i> Autophagy. <i>Brazilian Journal of Veterinary Parasitology</i> , 2021, 30, e017020.	0.7	2
31	Matrix metalloproteinase-14 regulates collagen degradation and migration of mononuclear cells during infection with genotype VII Newcastle disease virus. <i>Journal of General Virology</i> , 2021, 102, .	2.9	0
32	G1-like M and PB2 genes are preferentially incorporated into H7N9 progeny virions during genetic reassortment. <i>BMC Veterinary Research</i> , 2021, 17, 80.	1.9	0
33	Packaging signal of influenza A virus. <i>Virology Journal</i> , 2021, 18, 36.	3.4	27
34	Surveillance of Class I Newcastle Disease Virus at Live Bird Markets and Commercial Poultry Farms in Eastern China Reveals the Epidemic Characteristics. <i>Virologica Sinica</i> , 2021, 36, 818-822.	3.0	5
35	Mutations during the adaptation of H7N9 avian influenza virus to mice lungs enhance human-like sialic acid binding activity and virulence in mice. <i>Veterinary Microbiology</i> , 2021, 254, 109000.	1.9	4
36	Assay of extracellular matrix degradation and transmigration of chicken peripheral blood mononuclear cells after infection with genotype VII Newcastle disease virus in vitro. <i>Journal of Virological Methods</i> , 2021, 290, 114076.	2.1	0

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37	The Packaging Regions of G1-Like PB2 Gene Contribute to Improving the Survival Advantage of Genotype S H9N2 Virus in China. <i>Frontiers in Microbiology</i> , 2021, 12, 655057.	3.5	1
38	The virulence modulator PA-X protein has minor effect on the pathogenicity of the highly pathogenic H7N9 avian influenza virus in mice. <i>Veterinary Microbiology</i> , 2021, 255, 109019.	1.9	1
39	Identification of the dominant non-neutralizing epitope in the haemagglutinin of H7N9 avian influenza virus. <i>Virus Research</i> , 2021, 298, 198409.	2.2	4
40	Differential microRNA Expression in Newcastle Disease Virus-Infected HeLa Cells and Its Role in Regulating Virus Replication. <i>Frontiers in Oncology</i> , 2021, 11, 616809.	2.8	6
41	Electrospun Membranes as a Porous Barrier for Molecular Transport: Membrane Characterization and Release Assessment. <i>Pharmaceutics</i> , 2021, 13, 916.	4.5	6
42	Rapid differential detection of subtype H1 and H3 swine influenza viruses using a TaqMan-MGB-based duplex one-step real-time RT-PCR assay. <i>Archives of Virology</i> , 2021, 166, 2217-2224.	2.1	3
43	H7N9 influenza virus-like particle based on BEVS protects chickens from lethal challenge with highly pathogenic H7N9 avian influenza virus. <i>Veterinary Microbiology</i> , 2021, 258, 109106.	1.9	8
44	Antigen Camouflage and Decoy Strategy to Overcome Interference From Maternally Derived Antibody With Newcastle Disease Virus-Vectored Vaccines: More Than a Simple Combination. <i>Frontiers in Microbiology</i> , 2021, 12, 735250.	3.5	1
45	Genesis, evolution and host species distribution of influenza A (H10N3) virus in China. <i>Journal of Infection</i> , 2021, 83, 607-635.	3.3	7
46	AlphaB-crystallin promotes porcine circovirus type 2 replication in a cell proliferation-dependent manner. <i>Virus Research</i> , 2021, 301, 198435.	2.2	3
47	Genetic and antigenic diversity of H7N9 highly pathogenic avian influenza virus in China. <i>Infection, Genetics and Evolution</i> , 2021, 93, 104993.	2.3	12
48	G1-like PB2 gene improves virus replication and competitive advantage of H9N2 virus. <i>Virus Genes</i> , 2021, 57, 521-528.	1.6	1
49	Development of an Inactivated H7N9 Subtype Avian Influenza Serological DIVA Vaccine Using the Chimeric HA Epitope Approach. <i>Microbiology Spectrum</i> , 2021, 9, e0068721.	3.0	6
50	Common occurrence of <i>Enterocytozoon bieneusi</i> genotypes SHR1 and PL2 in farmed masked palm civet (<i>Paguma larvata</i>) in China. <i>International Journal for Parasitology: Parasites and Wildlife</i> , 2021, 16, 99-102.	1.5	1
51	A reassortant highly pathogenic avian influenza H5N6 virus originating from the wildbird-origin H5N6 and the poultry H9N2/H7N9 viruses in Xinjiang, China. <i>Medycyna Weterynaryjna</i> , 2021, 77, 6532-2021.	0.1	2
52	N-linked glycosylation at site 158 of the HA protein of H5N6 highly pathogenic avian influenza virus is important for viral biological properties and host immune responses. <i>Veterinary Research</i> , 2021, 52, 8.	3.0	19
53	Long noncoding RNA#45 exerts broad inhibitory effect on influenza A virus replication via its stem ring arms. <i>Virulence</i> , 2021, 12, 2443-2460.	4.4	7
54	Identification of a universal antigen epitope of influenza A virus using peptide microarray. <i>BMC Veterinary Research</i> , 2021, 17, 22.	1.9	1

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55	Biological Characterization and Evolutionary Dynamics of Pigeon Paramyxovirus Type 1 in China. <i>Frontiers in Veterinary Science</i> , 2021, 8, 721102.	2.2	2
56	gga-miR-1603 and gga-miR-1794 directly target viral L gene and function as a broad-spectrum antiviral factor against NDV replication. <i>Virulence</i> , 2021, 12, 45-56.	4.4	6
57	Experimental induction of necrotic enteritis with or without predisposing factors using netB positive <i>Clostridium perfringens</i> strains. <i>Gut Pathogens</i> , 2021, 13, 68.	3.4	6
58	Single Dose of Bivalent H5 and H7 Influenza Virus-Like Particle Protects Chickens Against Highly Pathogenic H5N1 and H7N9 Avian Influenza Viruses. <i>Frontiers in Veterinary Science</i> , 2021, 8, 774630.	2.2	6
59	Genome-Wide Analysis of Alternative Splicing during Host-Virus Interactions in Chicken. <i>Viruses</i> , 2021, 13, 2409.	3.3	3
60	Spatiotemporal Associations and Molecular Evolution of Highly Pathogenic Avian Influenza A H7N9 Virus in China from 2017 to 2021. <i>Viruses</i> , 2021, 13, 2524.	3.3	5
61	Rapid Emergence of the Reassortant 2.3.4.4b H5N2 Highly Pathogenic Avian Influenza Viruses in a Live Poultry Market in Xinjiang, Northwest China. <i>Avian Diseases</i> , 2021, 65, 578-583.	1.0	1
62	Characterization and evolution of the coronavirus porcine epidemic diarrhoea virus HLJBY isolated in China. <i>Transboundary and Emerging Diseases</i> , 2020, 67, 65-79.	3.0	20
63	Efficacy of the Bartha-K61 vaccine and a gE ^Δ /gI ^Δ /TK ^Δ prototype vaccine against variant porcine pseudorabies virus (vPRV) in piglets with sublethal challenge of vPRV. <i>Research in Veterinary Science</i> , 2020, 128, 16-23.	1.9	24
64	The PB2 and M genes are critical for the superiority of genotype S H9N2 virus to genotype H in optimizing viral fitness of H5Nx and H7N9 avian influenza viruses in mice. <i>Transboundary and Emerging Diseases</i> , 2020, 67, 758-768.	3.0	9
65	Amino acid substitutions in antigenic region B of hemagglutinin play a critical role in the antigenic drift of subclade 2.3.4.4 highly pathogenic H5NX influenza viruses. <i>Transboundary and Emerging Diseases</i> , 2020, 67, 263-275.	3.0	9
66	Occurrence and genotypes of <i>Cryptosporidium</i> spp., <i>Giardia duodenalis</i> , and <i>Blastocystis</i> sp. in household, shelter, breeding, and pet market dogs in Guangzhou, southern China. <i>Scientific Reports</i> , 2020, 10, 17736.	3.3	16
67	PA-X protein of H5N1 avian influenza virus inhibits NF-kappaB activity, a potential mechanism for PA-X counteracting the host innate immune responses. <i>Veterinary Microbiology</i> , 2020, 250, 108838.	1.9	5
68	Effect of different floatation solutions on <i>E. tenella</i> oocyst purification and optimization of centrifugation conditions for improved recovery of oocysts and sporocysts. <i>Experimental Parasitology</i> , 2020, 217, 107965.	1.2	2
69	Comparative pathogenicity of two closely related Newcastle disease virus isolates from chicken and pigeon respectively. <i>Virus Research</i> , 2020, 286, 198091.	2.2	10
70	Dominant subtype switch in avian influenza viruses during 2016–2019 in China. <i>Nature Communications</i> , 2020, 11, 5909.	12.8	93
71	A77 1726, the active metabolite of the anti-rheumatoid arthritis drug leflunomide, inhibits influenza A virus replication in vitro and in vivo by inhibiting the activity of Janus kinases. <i>FASEB Journal</i> , 2020, 34, 10132-10145.	0.5	15
72	H1N1 Influenza Virus Cross-Activates Gli1 to Disrupt the Intercellular Junctions of Alveolar Epithelial Cells. <i>Cell Reports</i> , 2020, 31, 107801.	6.4	28

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73	Truncation or Deglycosylation of the Neuraminidase Stalk Enhances the Pathogenicity of the H5N1 Subtype Avian Influenza Virus in Mallard Ducks. <i>Frontiers in Microbiology</i> , 2020, 11, 583588.	3.5	5
74	Detection of PB2 627K mutation in two highly pathogenic isolates of the H7N9 subtype Influenza A virus from chickens in Northern China. <i>Journal of Infection</i> , 2020, 81, 979-997.	3.3	5
75	Role of Post-translational Modifications in Influenza A Virus Life Cycle and Host Innate Immune Response. <i>Frontiers in Microbiology</i> , 2020, 11, 517461.	3.5	26
76	Establishing a Multicolor Flow Cytometry to Characterize Cellular Immune Response in Chickens Following H7N9 Avian Influenza Virus Infection. <i>Viruses</i> , 2020, 12, 1396.	3.3	17
77	Autophagy induced by monensin serves as a mechanism for programmed death in <i>Eimeria tenella</i> . <i>Veterinary Parasitology</i> , 2020, 287, 109181.	1.8	4
78	Substitutions in the PB2 methionine 283 residue affect H5 subtype avian influenza virus virulence. <i>Transboundary and Emerging Diseases</i> , 2020, 67, 2554-2563.	3.0	4
79	Newcastle Disease Virus as a Vaccine Vector for 20 Years: A Focus on Maternally Derived Antibody Interference. <i>Vaccines</i> , 2020, 8, 222.	4.4	29
80	Colonisation of mice and pigs by a chimeric porcine circovirus 1 prototype vaccine strain and a PCV2 isolate originating in China and their induction of cytokines. <i>Journal of Virological Methods</i> , 2020, 283, 113905.	2.1	1
81	<i>Cryptosporidium parvum</i> gp40/15 Is Associated with the Parasitophorous Vacuole Membrane and Is a Potential Vaccine Target. <i>Microorganisms</i> , 2020, 8, 363.	3.6	11
82	Induction of cross-group broadly reactive antibody response by natural H7N9 avian influenza virus infection and immunization with inactivated H7N9 vaccine in chickens. <i>Transboundary and Emerging Diseases</i> , 2020, 67, 3041-3048.	3.0	2
83	Inhibition of porcine epidemic diarrhea virus (PEDV) replication by A77 1726 through targeting JAK and Src tyrosine kinases. <i>Virology</i> , 2020, 551, 75-83.	2.4	11
84	Pathogenicity and transmissibility of clade 2.3.4.4 highly pathogenic avian influenza virus subtype H5N6 in pigeons. <i>Veterinary Microbiology</i> , 2020, 247, 108776.	1.9	4
85	Glycosylation deletion of hemagglutinin head in the H5 subtype avian influenza virus enhances its virulence in mammals by inducing endoplasmic reticulum stress. <i>Transboundary and Emerging Diseases</i> , 2020, 67, 1492-1506.	3.0	7
86	Characterization and functional analysis of chicken APOBEC4. <i>Developmental and Comparative Immunology</i> , 2020, 106, 103631.	2.3	9
87	Isolation and characterization of Getah virus from pigs in Guangdong province of China. <i>Transboundary and Emerging Diseases</i> , 2020, 67, 2249.	3.0	23
88	EntE, EntS and TolC synergistically contributed to the pathogenesis of APEC strain E058. <i>Microbial Pathogenesis</i> , 2020, 141, 103990.	2.9	9
89	Effect of the selection pressure of vaccine antibodies on evolution of H9N2 avian influenza virus in chickens. <i>AMB Express</i> , 2020, 10, 98.	3.0	16
90	Role of the Hemagglutinin Residue 227 in Immunogenicity of H5 and H7 Subtype Avian Influenza Vaccines in Chickens. <i>Avian Diseases</i> , 2020, 64, 445-450.	1.0	1

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91	MicroRNA Expression Profiling in Newcastle Disease Virus-Infected DF-1 Cells by Deep Sequencing. <i>Frontiers in Microbiology</i> , 2019, 10, 1659.	3.5	15
92	The PB2 and M genes of genotype S H9N2 virus contribute to the enhanced fitness of H5Nx and H7N9 avian influenza viruses in chickens. <i>Virology</i> , 2019, 535, 218-226.	2.4	13
93	Sensorless-Based Active Disturbance Rejection Control for a Wind Energy Conversion System With Permanent Magnet Synchronous Generator. <i>IEEE Access</i> , 2019, 7, 122663-122674.	4.2	28
94	Gga-miR-19b-3p Inhibits Newcastle Disease Virus Replication by Suppressing Inflammatory Response via Targeting RNF11 and ZMYND11. <i>Frontiers in Microbiology</i> , 2019, 10, 2006.	3.5	17
95	Role of TGF- β -activated kinase 1 (TAK1) activation in H5N1 influenza A virus-induced c-Jun terminal kinase activation and virus replication. <i>Virology</i> , 2019, 537, 263-271.	2.4	8
96	Recombinant baculovirus vaccine expressing hemagglutinin of H7N9 avian influenza virus confers full protection against lethal highly pathogenic H7N9 virus infection in chickens. <i>Archives of Virology</i> , 2019, 164, 807-817.	2.1	8
97	Non-linear extended state observer-based sliding mode control for a direct-driven wind energy conversion system with permanent magnet synchronous generator. <i>Journal of Engineering</i> , 2019, 2019, 613-617.	1.1	9
98	The effect of autophagy on the survival and invasive activity of <i>Eimeria tenella</i> sporozoites. <i>Scientific Reports</i> , 2019, 9, 5835.	3.3	7
99	Multiplex one-step real-time PCR assay for rapid simultaneous detection of velogenic and mesogenic Newcastle disease virus and H5-subtype avian influenza virus. <i>Archives of Virology</i> , 2019, 164, 1111-1119.	2.1	7
100	Enhanced cross-lineage protection induced by recombinant H9N2 avian influenza virus inactivated vaccine. <i>Vaccine</i> , 2019, 37, 1736-1742.	3.8	9
101	Compound control method for DC-DC converter. <i>Journal of Engineering</i> , 2019, 2019, 8348-8352.	1.1	1
102	Catalytic inactivation of influenza virus by iron oxide nanozyme. <i>Theranostics</i> , 2019, 9, 6920-6935.	10.0	90
103	Speed sensorless model predictive control method for a direct-drive wind energy conversion system. <i>Measurement and Control</i> , 2019, 52, 1394-1402.	1.8	5
104	Impact of the variations in potential glycosylation sites of the hemagglutinin of H9N2 influenza virus. <i>Virus Genes</i> , 2019, 55, 182-190.	1.6	11
105	Unexpected transcriptome <i>pompT</i> ¹ contributes to the increased pathogenicity of a <i>pompT</i> mutant of avian pathogenic <i>Escherichia coli</i> . <i>Veterinary Microbiology</i> , 2019, 228, 61-68.	1.9	2
106	The virulence of NDV NA-1 strain regulated by the 3' leader or 5' trailer sequences. <i>Microbial Pathogenesis</i> , 2019, 126, 109-115.	2.9	0
107	Role of c-Jun terminal kinase (JNK) activation in influenza A virus-induced autophagy and replication. <i>Virology</i> , 2019, 526, 1-12.	2.4	37
108	Hemagglutinin-Specific Non-neutralizing Antibody Is Essential for Protection Provided by Inactivated and Viral-Vectored H7N9 Avian Influenza Vaccines in Chickens. <i>Frontiers in Veterinary Science</i> , 2019, 6, 482.	2.2	12

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109	Redescription of <i>Bryobia pritchardi</i> Rimando, 1962 (Acari: Tetranychidae), with an ontogeny of chaetotaxy. <i>Acarologia</i> , 2019, 59, 73-110.	0.6	0
110	T160A mutation-induced deglycosylation at site 158 in hemagglutinin is a critical determinant of the dual receptor binding properties of clade 2.3.4.4 H5NX subtype avian influenza viruses. <i>Veterinary Microbiology</i> , 2018, 217, 158-166.	1.9	25
111	Genetic and biological characterization of three poultry-origin H5N6 avian influenza viruses with all internal genes from genotype S H9N2 viruses. <i>Archives of Virology</i> , 2018, 163, 947-960.	2.1	12
112	Quantitative proteomics identify an association between extracellular matrix degradation and immunopathology of genotype VII Newcastle disease virus in the spleen in chickens. <i>Journal of Proteomics</i> , 2018, 181, 201-212.	2.4	13
113	Genetic analysis and biological characteristics of different internal gene origin H5N6 reassortment avian influenza virus in China in 2016. <i>Veterinary Microbiology</i> , 2018, 219, 200-211.	1.9	19
114	NDV entry into dendritic cells through macropinocytosis and suppression of T lymphocyte proliferation. <i>Virology</i> , 2018, 518, 126-135.	2.4	12
115	Evolution of H9N2 avian influenza virus in embryonated chicken eggs with or without homologous vaccine antibodies. <i>BMC Veterinary Research</i> , 2018, 14, 71.	1.9	12
116	New Threats from H7N9 Influenza Virus: Spread and Evolution of High- and Low-Pathogenicity Variants with High Genomic Diversity in Wave Five. <i>Journal of Virology</i> , 2018, 92, .	3.4	92
117	Importin $\beta 5$ negatively regulates importin $\beta 1$ -mediated nuclear import of Newcastle disease virus matrix protein and viral replication and pathogenicity in chicken fibroblasts. <i>Virulence</i> , 2018, 9, 783-803.	4.4	23
118	Characteristics of the emerging chicken-origin highly pathogenic H7N9 viruses: A new threat to public health and poultry industry. <i>Journal of Infection</i> , 2018, 76, 217-220.	3.3	29
119	Composite Anti-Disturbance Control of Permanent Magnet Synchronous Motor Based on Feedback Linearization. , 2018, , .		0
120	Design of an Intelligent Active Obstacle Avoidance Car Based on Rotating Ultrasonic Sensors. , 2018, , .		13
121	An RBFNN-Based Direct Inverse Controller for PMSM with Disturbances. <i>Complexity</i> , 2018, 2018, 1-13.	1.6	1
122	A Model-Compensation ADRC Strategy of Wind Energy Conversion System with Direct-Driven PMSG. , 2018, , .		2
123	Ontogenetic development and redescription of <i>Eotetranychus kankitus</i> (Acariformes: Tetranychidae). <i>Zootaxa</i> , 2018, 4540, 132.	0.5	1
124	Two novel reassortant high pathogenic H7N9 viruses isolated in Southern China in fifth wave shows internal genomic diversity and high virulence in chickens and ducks. <i>Journal of Infection</i> , 2018, 77, 561-571.	3.3	4
125	Chimeric Newcastle disease virus-vectored vaccine protects chickens against H9N2 avian influenza virus in the presence of pre-existing NDV immunity. <i>Archives of Virology</i> , 2018, 163, 3365-3371.	2.1	15
126	Development of a Colloidal Gold-Based Immunochromatographic Strip for Rapid Detection of H7N9 Influenza Viruses. <i>Frontiers in Microbiology</i> , 2018, 9, 2069.	3.5	21

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127	Down-Regulation of <i>SSII-2</i> Gene Expression Results in Novel Low-Amylose Rice with Soft, Transparent Grains. <i>Journal of Agricultural and Food Chemistry</i> , 2018, 66, 9750-9760.	5.2	28
128	Retrospective survey and phylogenetic analysis of porcine circovirus type 3 in Jiangsu province, China, 2008 to 2017. <i>Archives of Virology</i> , 2018, 163, 2531-2538.	2.1	23
129	Signature-tagged mutagenesis screening revealed the role of lipopolysaccharide biosynthesis gene <i>rfbH</i> in smooth-to-rough transition in <i>Salmonella Enteritidis</i> . <i>Microbiological Research</i> , 2018, 212-213, 75-79.	5.3	6
130	Genetic and biological characterization of H9N2 avian influenza viruses isolated in China from 2011 to 2014. <i>PLoS ONE</i> , 2018, 13, e0199260.	2.5	23
131	PA-X: a key regulator of influenza A virus pathogenicity and host immune responses. <i>Medical Microbiology and Immunology</i> , 2018, 207, 255-269.	4.8	32
132	Deep sequencing of the mouse lung transcriptome reveals distinct long non-coding RNAs expression associated with the high virulence of H5N1 avian influenza virus in mice. <i>Virulence</i> , 2018, 9, 1092-1111.	4.4	7
133	Characterization of cattle-origin ticks from Southern China. <i>Acta Tropica</i> , 2018, 187, 92-98.	2.0	9
134	Effect of annexin II-mediated conversion of plasmin from plasminogen on airborne transmission of H9N2 avian influenza virus. <i>Veterinary Microbiology</i> , 2018, 223, 100-106.	1.9	11
135	Evaluation of the Efficacy and Cross-Protective Immunity of Live-Attenuated Chimeric PCV1-2b Vaccine Against PCV2b and PCV2d Subtype Challenge in Pigs. <i>Frontiers in Microbiology</i> , 2018, 9, 455.	3.5	16
136	The PA-interacting host protein nucleolin acts as an antiviral factor during highly pathogenic H5N1 avian influenza virus infection. <i>Archives of Virology</i> , 2018, 163, 2775-2786.	2.1	10
137	A comprehensive comparison of the fifth wave highly pathogenic and low pathogenic H7N9 avian influenza viruses reveals potential threat posed by both types of viruses in mammals. <i>Transboundary and Emerging Diseases</i> , 2018, 65, 1459-1473.	3.0	10
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