Zuzhang Wei

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	High prevalence of a novel porcine bocavirus in weanling piglets with respiratory tract symptoms in China. Archives of Virology, 2010, 155, 1313-1317.	0.9	83
2	Arterivirus Minor Envelope Proteins Are a Major Determinant of Viral Tropism in Cell Culture. Journal of Virology, 2012, 86, 3701-3712.	1.5	78
3	Host miR-26a suppresses replication of porcine reproductive and respiratory syndrome virus by upregulating type I interferons. Virus Research, 2015, 195, 86-94.	1.1	71
4	N-Linked Glycosylation of GP5 of Porcine Reproductive and Respiratory Syndrome Virus Is Critically Important for Virus Replication <i>In Vivo</i> . Journal of Virology, 2012, 86, 9941-9951.	1.5	60
5	Emergence and Evolution of Novel Reassortant Influenza A Viruses in Canines in Southern China. MBio, 2018, 9, .	1.8	41
6	Construction of infectious cDNA clones of PRRSV: Separation of coding regions for nonstructural and structural proteins. Science in China Series C: Life Sciences, 2008, 51, 271-279.	1.3	38
7	Porcine reproductive and respiratory syndrome virus ORF5a protein is essential for virus viability. Virus Research, 2013, 171, 178-185.	1.1	31
8	Novel triple-reassortant influenza viruses in pigs, Guangxi, China. Emerging Microbes and Infections, 2018, 7, 1-9.	3.0	31
9	A 5'-proximal Stem-loop Structure of 5' Untranslated Region of Porcine Reproductive and Respiratory Syndrome Virus Genome Is Key for Virus Replication. Virology Journal, 2011, 8, 172.	1.4	26
10	Pathogenic Characteristics of a Porcine Astrovirus Strain Isolated in China. Viruses, 2019, 11, 1156.	1.5	26
11	Influence of N-linked glycosylation of minor proteins of porcine reproductive and respiratory syndrome virus on infectious virus recovery and receptor interaction. Virology, 2012, 429, 1-11.	1.1	25
12	Immunization of pigs with a type 2 modified live PRRSV vaccine prevents the development of a deadly long lasting hyperpyrexia in a challenge study with highly pathogenic PRRSV JX143. Vaccine, 2013, 31, 2062-2066.	1.7	24
13	Genetic analysis of porcine circovirus type 2 (PCV2) strains between 2002 and 2016 reveals PCV2 mutant predominating in porcine population in Guangxi, China. BMC Veterinary Research, 2019, 15, 118.	0.7	21
14	Cis-acting structural element in 5′ UTR is essential for infectivity of porcine reproductive and respiratory syndrome virus. Virus Research, 2012, 163, 108-119.	1.1	20
15	Molecular epidemiology and viremia of porcine astrovirus in pigs from Guangxi province of China. BMC Veterinary Research, 2019, 15, 471.	0.7	20
16	Construction of a reverse genetic system for porcine astrovirus. Archives of Virology, 2018, 163, 1511-1518.	0.9	19
17	Detection and genetic characterization of canine astroviruses in pet dogs in Guangxi, China. Virology Journal, 2017, 14, 156.	1.4	18
18	Development of a differentiable virus via a spontaneous deletion in the nsp2 region associated with cell adaptation of porcine reproductive and respiratory syndrome virus. Virus Research, 2013, 171, 150-160.	1.1	17

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19	Activation of RNase L in Egyptian Rousette Bat-Derived RoNi/7 Cells Is Dependent Primarily on OAS3 and Independent of MAVS Signaling. MBio, 2019, 10, .	1.8	17
20	Genetic Diversity of Porcine Epidemic Diarrhea Virus With a Naturally Occurring Truncated ORF3 Gene Found in Guangxi, China. Frontiers in Veterinary Science, 2020, 7, 435.	0.9	14
21	Emergence and Phylogenetic Analysis of a Getah Virus Isolated in Southern China. Frontiers in Veterinary Science, 2020, 7, 552517.	0.9	14
22	Characterization of swine-origin H1N1 canine influenza viruses. Emerging Microbes and Infections, 2019, 8, 1017-1026.	3.0	13
23	Generation of a porcine reproductive and respiratory syndrome virus expressing a marker gene inserted between ORF4 and ORF5a. Archives of Virology, 2020, 165, 1803-1813.	0.9	13
24	Comparative Characterization and Pathogenicity of a Novel Porcine Epidemic Diarrhea Virus (PEDV) with a Naturally Occurring Truncated ORF3 Gene Coinfected with PEDVs Possessing an Intact ORF3 Gene in Piglets. Viruses, 2021, 13, 1562.	1.5	13
25	Identification of non-essential regions in nucleocapsid protein of porcine reproductive and respiratory syndrome virus for replication in cell culture. Virus Research, 2011, 158, 62-71.	1.1	12
26	Genetic analysis of porcine productive and respiratory syndrome virus between 2013 and 2014 in Southern parts of China: identification of several novel strains with amino acid deletions or insertions in nsp2. BMC Veterinary Research, 2019, 15, 171.	0.7	11
27	Full Genomic Analysis of New Variants of Porcine Reproductive and Respiratory Syndrome Virus Revealed Multiple Recombination Events Between Different Lineages and Sublineages. Frontiers in Veterinary Science, 2020, 7, 603.	0.9	11
28	Replacement of the heterologous 5′ untranslated region allows preservation of the fully functional activities of type 2 porcine reproductive and respiratory syndrome virus. Virology, 2013, 439, 1-12.	1.1	10
29	Emergence and phylogenetic analysis of a novel Seneca Valley virus strain in the Guangxi Province of China. Research in Veterinary Science, 2020, 130, 207-211.	0.9	10
30	Coâ€circulation and evolution of genogroups I and II of respiratory and enteric feline calicivirus isolates in cats. Transboundary and Emerging Diseases, 2022, 69, 2924-2937.	1.3	10
31	Genetic manipulation of a transcription-regulating sequence of porcine reproductive and respiratory syndrome virus reveals key nucleotides determining its activity. Archives of Virology, 2014, 159, 1927-1940.	0.9	8
32	Effect of an 88-amino-acid deletion in nsp2 of porcine reproductive and respiratory syndrome virus on virus replication and cytokine responses in vitro. Archives of Virology, 2018, 163, 1489-1501.	0.9	8
33	Construction and characterization of a full-length infectious clone of Getah virus in vivo. Virologica Sinica, 2022, 37, 348-357.	1.2	8
34	Establishment of a Multiplex RT-PCR Method for the Detection of Five Known Genotypes of Porcine Astroviruses. Frontiers in Veterinary Science, 2021, 8, 684279.	0.9	7
35	Isolation, Identification, and Evaluation of the Pathogenicity of a Porcine Enterovirus G Isolated From China. Frontiers in Veterinary Science, 2021, 8, 712679.	0.9	7
36	Conserved nucleotides in the terminus of the 3′ UTR region are important for the replication and infectivity of porcine reproductive and respiratory syndrome virus. Archives of Virology, 2013, 158, 1719-1732.	0.9	6

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37	Cysteine residues of the porcine reproductive and respiratory syndrome virus ORF5a protein are not essential for virus viability. Virus Research, 2015, 197, 17-25.	1.1	5
38	Generation of a Recombinant Porcine Reproductive and Respiratory Syndrome Virus Stably Expressing Two Marker Genes. Frontiers in Veterinary Science, 2020, 7, 548282.	0.9	5
39	Identification of a novel protein in porcine astrovirus that is important for virus replication. Veterinary Microbiology, 2021, 255, 108984.	0.8	5
40	Genetic Characteristics and Pathogenicity of a Novel Porcine Deltacoronavirus Southeast Asia-Like Strain Found in China. Frontiers in Veterinary Science, 2021, 8, 701612.	0.9	5
41	Complete Genome Sequence of Feline Calicivirus Strain GX01-2013 Isolated from Household Cats in Guangxi, Southern China. Genome Announcements, 2016, 4, .	0.8	4
42	Phylogenetic Diversity of Classical Swine Fever Virus (CSFV) Field Isolates from Outbreaks in China Between 2008 and 2011. Asian Journal of Animal and Veterinary Advances, 2013, 8, 449-460.	0.3	4
43	Use of reverse genetics to develop a novel marker porcine reproductive and respiratory syndrome virus. Virus Genes, 2012, 45, 548-555.	0.7	3
44	Identification of novel B-cell epitopes on the capsid protein of type 1 porcine astrovirus, using monoclonal antibodies. International Journal of Biological Macromolecules, 2021, 189, 939-947.	3.6	2
45	Insertion of Exogenous Genes within the ORF1a Coding Region of Porcine Astrovirus. Viruses, 2021, 13, 2119.	1.5	2
46	Detection and Genetic Diversity of a Novel Water Buffalo Astrovirus Species Found in the Guangxi Province of China. Frontiers in Veterinary Science, 2021, 8, 692193.	0.9	1
47	Isolation and Phylogenetic Analysis of a Hunnivirus Strain in Water Buffaloes From China. Frontiers in Veterinary Science. 2022. 9. 851743.	0.9	1