

Shao-Lun Zhai

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8720888/publications.pdf>

Version: 2024-02-01

33
papers

552
citations

687363

13
h-index

642732

23
g-index

35
all docs

35
docs citations

35
times ranked

801
citing authors

#	ARTICLE	IF	CITATIONS
1	Comparative epidemiology of porcine circovirus type 3 in pigs with different clinical presentations. <i>Virology Journal</i> , 2017, 14, 222.	3.4	75
2	Porcine circovirus type 2 in China: an update on and insights to its prevalence and control. <i>Virology Journal</i> , 2014, 11, 88.	3.4	74
3	High prevalence of bovine viral diarrhoea virus 1 in Chinese swine herds. <i>Veterinary Microbiology</i> , 2012, 159, 490-493.	1.9	65
4	Reservoirs of Porcine Circoviruses: A Mini Review. <i>Frontiers in Veterinary Science</i> , 2019, 6, 319.	2.2	45
5	Occurrence and sequence analysis of porcine deltacoronaviruses in southern China. <i>Virology Journal</i> , 2016, 13, 136.	3.4	27
6	Genetic Variability and Phylogeny of Current Chinese Porcine Epidemic Diarrhoea Virus Strains Based on Spike, ORF3, and Membrane Genes. <i>Scientific World Journal</i> , The, 2014, 2014, 1-8.	2.1	21
7	Emergence of reticuloendotheliosis virus in pigeons in Guangdong Province, Southern China. <i>Archives of Virology</i> , 2016, 161, 2007-2011.	2.1	20
8	Where did SARS-CoV-2 come from?. <i>Veterinary Record</i> , 2020, 186, 254-254.	0.3	20
9	Outbreak and genotyping of canine distemper virus in captive Siberian tigers and red pandas. <i>Scientific Reports</i> , 2017, 7, 8132.	3.3	19
10	African swine fever spread in China. <i>Veterinary Record</i> , 2019, 184, 559-559.	0.3	19
11	First complete genome sequence of parainfluenza virus 5 isolated from lesser panda. <i>Archives of Virology</i> , 2017, 162, 1413-1418.	2.1	18
12	Molecular detection and genome characterization of porcine circovirus type 2 in rats captured on commercial swine farms. <i>Archives of Virology</i> , 2016, 161, 3237-3244.	2.1	15
13	Development and Characterization of a Reverse-Genetics System for Influenza D Virus. <i>Journal of Virology</i> , 2019, 93, .	3.4	15
14	Complete genome characterization and phylogenetic analysis of three distinct buffalo-origin PCV2 isolates from China. <i>Infection, Genetics and Evolution</i> , 2014, 28, 278-282.	2.3	12
15	Preliminary serological evidence for Schmallenberg virus infection in China. <i>Tropical Animal Health and Production</i> , 2018, 50, 449-453.	1.4	12
16	Phylogeographic analysis of porcine reproductive and respiratory syndrome virus 1 in Guangdong province, Southern China. <i>Archives of Virology</i> , 2018, 163, 2443-2449.	2.1	11
17	A novel porcine kobuvirus emerged in piglets with severe diarrhoea in China. <i>Transboundary and Emerging Diseases</i> , 2017, 64, 1030-1036.	3.0	10
18	Genome characterization and phylogenetic analysis of a lineage IV peste des petits ruminants virus in southern China. <i>Virus Genes</i> , 2015, 51, 361-366.	1.6	9

#	ARTICLE	IF	CITATIONS
19	High prevalence of torque teno sus virus in China and genetic diversity of the 5' non-coding region. Archives of Virology, 2013, 158, 1567-1573.	2.1	8
20	Phylogeography and evolutionary dynamics analysis of porcine delta coronavirus with host expansion to humans. Transboundary and Emerging Diseases, 2022, 69, .	3.0	8
21	Buffalo-Origin Seneca Valley Virus in China: First Report, Isolation, Genome Characterization, and Evolution Analysis. Frontiers in Veterinary Science, 2021, 8, 730701.	2.2	7
22	First molecular detection of porcine circovirus type 2 in bovids in China. Virus Genes, 2014, 49, 507-511.	1.6	6
23	Reappearance of buffalo-origin-like porcine circovirus type 2 strains in swine herds in southern China. New Microbes and New Infections, 2017, 17, 98-100.	1.6	6
24	A Novel Enzootic Nasal Tumor Virus Circulating in Goats from Southern China. Viruses, 2019, 11, 956.	3.3	5
25	First complete genome sequence of canine bocavirus 2 in mainland China. New Microbes and New Infections, 2017, 18, 47-49.	1.6	4
26	Natural infections of SARS-CoV-2 increased in animals: How should humans interact with animals?. Journal of Medical Virology, 2022, 94, 3503-3505.	5.0	4
27	Can cats become infected with Covid-19?. Veterinary Record, 2020, 186, e20.	0.3	3
28	Threat of lumpy skin disease to the Chinese cattle industry. Veterinary Record, 2021, 188, 315-316.	0.3	3
29	Development of a Real-Time Quantitative RT-PCR Assay for Detection of Bovine Rhinitis B Virus. Frontiers in Veterinary Science, 2021, 8, 680707.	2.2	3
30	Diverse african swine fever viruses in China. New Microbes and New Infections, 2022, 46, 100976.	1.6	3
31	Ruminal impaction due to Ficus esquiroliana Levl. in Boer goats. SpringerPlus, 2013, 2, 608.	1.2	2
32	Genome Characterization and Phylogenetic Analysis of the First Bovine Rhinitis B Virus Isolate in China. Frontiers in Veterinary Science, 2021, 8, 721284.	2.2	2
33	Can porcine circovirus type 3 cause persistent infection in pigs?. Veterinary Record, 2019, 184, 617-618.	0.3	0