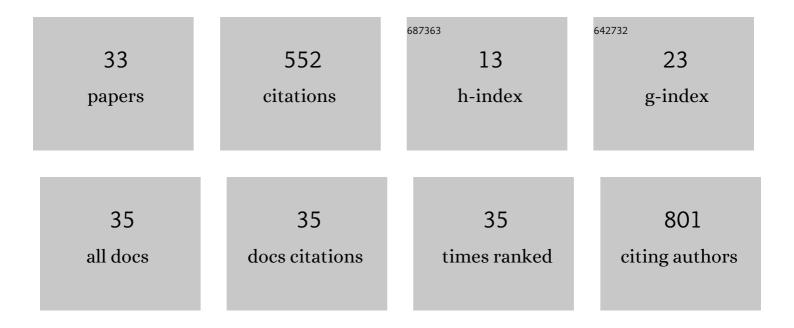
## Shao-Lun Zhai

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8720888/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Diverse african swine fever viruses in China. New Microbes and New Infections, 2022, 46, 100976.	1.6	3
2	Phylogeography and evolutionary dynamics analysis of porcine delta oronavirus with host expansion to humans. Transboundary and Emerging Diseases, 2022, 69, .	3.0	8
3	Natural infections of SARSâ€CoVâ€2 increased in animals: How should humans interactÂwith animals?. Journal of Medical Virology, 2022, 94, 3503-3505.	5.0	4
4	Threat of lumpy skin disease to the Chinese cattle industry. Veterinary Record, 2021, 188, 315-316.	0.3	3
5	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
6	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
7	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
8	Can cats become infected with Covidâ€19?. Veterinary Record, 2020, 186, e20.	0.3	3
9	Where did SARSâ€CoVâ€⊋ come from?. Veterinary Record, 2020, 186, 254-254.	0.3	20
10	Development and Characterization of a Reverse-Genetics System for Influenza D Virus. Journal of Virology, 2019, 93, .	3.4	15
11	Reservoirs of Porcine Circoviruses: A Mini Review. Frontiers in Veterinary Science, 2019, 6, 319.	2.2	45
12	A Novel Enzootic Nasal Tumor Virus Circulating in Goats from Southern China. Viruses, 2019, 11, 956.	3.3	5
13	Can porcine circovirus type 3 cause persistent infection in pigs?. Veterinary Record, 2019, 184, 617-618.	0.3	Ο
14	African swine fever spread in China. Veterinary Record, 2019, 184, 559-559.	0.3	19
15	Preliminary serological evidence for Schmallenberg virus infection in China. Tropical Animal Health and Production, 2018, 50, 449-453.	1.4	12
16	Phylogeographic analysis of porcine reproductive and respiratory syndrome virus 1 in Guangdong province, Southern China. Archives of Virology, 2018, 163, 2443-2449.	2.1	11
17	First complete genome sequence of parainfluenza virus 5 isolated from lesser panda. Archives of Virology, 2017, 162, 1413-1418.	2.1	18
18	A novel porcine kobuvirus emerged in piglets with severe diarrhoea in China. Transboundary and Emerging Diseases, 2017, 64, 1030-1036.	3.0	10

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#	Article	IF	CITATIONS
19	First complete genome sequence of canine bocavirus 2 in mainland China. New Microbes and New Infections, 2017, 18, 47-49.	1.6	4
20	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
21	Outbreak and genotyping of canine distemper virus in captive Siberian tigers and red pandas. Scientific Reports, 2017, 7, 8132.	3.3	19
22	Comparative epidemiology of porcine circovirus type 3 in pigs with different clinical presentations. Virology Journal, 2017, 14, 222.	3.4	75
23	Occurrence and sequence analysis of porcine deltacoronaviruses in southern China. Virology Journal, 2016, 13, 136.	3.4	27
24	Molecular detection and genome characterization of porcine circovirus type 2 in rats captured on commercial swine farms. Archives of Virology, 2016, 161, 3237-3244.	2.1	15
25	Emergence of reticuloendotheliosis virus in pigeons in Guangdong Province, Southern China. Archives of Virology, 2016, 161, 2007-2011.	2.1	20
26	Genome characterization and phylogenetic analysis of a lineage IV peste des petits ruminants virus in southern China. Virus Genes, 2015, 51, 361-366.	1.6	9
27	Genetic Variability and Phylogeny of Current Chinese Porcine Epidemic Diarrhea Virus Strains Based onSpike,ORF3, andMembraneGenes. Scientific World Journal, The, 2014, 2014, 1-8.	2.1	21
28	First molecular detection of porcine circovirus type 2 in bovids in China. Virus Genes, 2014, 49, 507-511.	1.6	6
29	Complete genome characterization and phylogenetic analysis of three distinct buffalo-origin PCV2 isolates from China. Infection, Genetics and Evolution, 2014, 28, 278-282.	2.3	12
30	Porcine circovirus type 2 in China: an update on and insights to its prevalence and control. Virology Journal, 2014, 11, 88.	3.4	74
31	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
32	Ruminal impaction due to Ficus esquiroliana Levl. in Boer goats. SpringerPlus, 2013, 2, 608.	1.2	2
33	High prevalence of bovine viral diarrhea virus 1 in Chinese swine herds. Veterinary Microbiology, 2012, 159, 490-493.	1.9	65