

Yan-Kuo Sun

List of Publications by Year in descending order

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16
papers

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1040056

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17
times ranked

316
citing authors

#	ARTICLE	IF	CITATIONS
1	First identification of porcine parvovirus 7 in China. Archives of Virology, 2018, 163, 209-213.	2.1	42
2	Inhibition of proanthocyanidin A2 on porcine reproductive and respiratory syndrome virus replication in vitro. PLoS ONE, 2018, 13, e0193309.	2.5	28
3	Platycodin D Suppresses Type 2 Porcine Reproductive and Respiratory Syndrome Virus In Primary and Established Cell Lines. Viruses, 2018, 10, 657.	3.3	23
4	First report and genetic characterization of feline kobuvirus in diarrhoeic cats in China. Transboundary and Emerging Diseases, 2018, 65, 1357-1363.	3.0	21
5	Phylogeography, phylodynamics and the recent outbreak of lineage 3 porcine reproductive and respiratory syndrome viruses in China. Transboundary and Emerging Diseases, 2019, 66, 2152-2162.	3.0	21
6	Insights into the evolutionary history and epidemiological characteristics of the emerging lineage 1 porcine reproductive and respiratory syndrome viruses in China. Transboundary and Emerging Diseases, 2020, 67, 2630-2641.	3.0	17
7	Emergence of novel recombination lineage 3 of porcine reproductive and respiratory syndrome viruses in Southern China. Transboundary and Emerging Diseases, 2019, 66, 578-587.	3.0	16
8	Genetic variation, pathogenicity, and immunogenicity of highly pathogenic porcine reproductive and respiratory syndrome virus strain XH-GD at different passage levels. Archives of Virology, 2016, 161, 77-86.	2.1	13
9	Isolation, identification and phylogenetic analysis of lumpy skin disease virus strain of outbreak in Guangdong, China. Transboundary and Emerging Diseases, 2022, 69, .	3.0	13
10	Porcine epidemic diarrhea virus in Asia: An alarming threat to the global pig industry. Infection, Genetics and Evolution, 2019, 70, 24-26.	2.3	12
11	Natural recombination of equine hepacivirus subtype 1 within the NS5A and NS5B genes. Virology, 2019, 533, 93-98.	2.4	8
12	Identification and genome characterization of a novel feline picornavirus proposed in the Hunnivirus genus. Infection, Genetics and Evolution, 2019, 71, 47-50.	2.3	7
13	First report and genetic diversity of porcine bufavirus in China. Virology Journal, 2020, 17, 2.	3.4	6
14	Identification and molecular analysis of Ixodid ticks (Acari: Ixodidae) infesting wild boars (Sus scrofa) and tick-borne pathogens at the Meihua mountain of southwestern Fujian, China. Veterinary Parasitology: Regional Studies and Reports, 2020, 22, 100492.	0.5	6
15	Lack of exposure of H10N8 avian influenza virus among veterinarians in guangdong province, China. Journal of Medical Virology, 2015, 87, 2018-2020.	5.0	3
16	Whole genome sequencing of clinical specimens reveals the genomic diversity of porcine reproductive and respiratory syndrome viruses emerging in China. Transboundary and Emerging Diseases, 2022, 69, .	3.0	2