

Cheng Tang

List of Publications by Year in descending order

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44
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#	ARTICLE	IF	CITATIONS
1	First Detection and Genomic Characterization of Bovine Norovirus from Yak. <i>Pathogens</i> , 2022, 11, 192.	2.8	3
2	Genomic analyses of <i>Staphylococcus aureus</i> isolated from yaks in Ganzi Tibetan Autonomous Prefecture, China. <i>Journal of Antimicrobial Chemotherapy</i> , 2022, , .	3.0	1
3	Aichivirus C isolate is a diarrhoea-causing pathogen in goats. <i>Transboundary and Emerging Diseases</i> , 2022, 69, .	3.0	4
4	Circulation of heterogeneous <i>Carnivore protoparvovirus 1</i> in diarrheal cats and prevalence of an A91S feline panleukopenia virus variant in China. <i>Transboundary and Emerging Diseases</i> , 2022, 69, .	3.0	5
5	Diversity of Astrovirus in Goats in Southwest China and Identification of Two Novel Caprine Astroviruses. <i>Microbiology Spectrum</i> , 2022, 10, .	3.0	2
6	Detection and molecular characteristics of bovine rotavirus A in dairy calves in China. <i>Journal of Veterinary Science</i> , 2021, 22, e69.	1.3	8
7	Genomic characterization of a nebovirus strain with a novel RdRp genotype in yaks. <i>Archives of Virology</i> , 2021, 166, 967-972.	2.1	0
8	Genomic analysis reveals G3P[13] porcine rotavirus A interspecific transmission to human from pigs in a swine farm with diarrhoea outbreak. <i>Journal of General Virology</i> , 2021, 102, .	2.9	3
9	Development and use of a reverse transcription insulated isothermal PCR assay for detection and characterization of bovine torovirus in yaks. <i>Archives of Virology</i> , 2021, 166, 2017-2025.	2.1	3
10	Identification of a novel Aichivirus D in sheep. <i>Infection, Genetics and Evolution</i> , 2021, 91, 104810.	2.3	8
11	Identification and genomic characterization of bovine parvovirus 1 in yaks. <i>Journal of Veterinary Diagnostic Investigation</i> , 2021, 33, 104063872110296.	1.1	0
12	First detection of canine norovirus in dogs and a complete GVI.2 genome in mainland China. <i>Infection, Genetics and Evolution</i> , 2021, 92, 104879.	2.3	3
13	Identification of a novel astrovirus in goats in China. <i>Infection, Genetics and Evolution</i> , 2021, 96, 105105.	2.3	5
14	First detection and molecular characteristics of caprine kobuvirus in goats in China. <i>Infection, Genetics and Evolution</i> , 2020, 85, 104566.	2.3	8
15	An emerging novel bovine coronavirus with a 4-amino-acid insertion in the receptor-binding domain of the hemagglutinin-esterase gene. <i>Archives of Virology</i> , 2020, 165, 3011-3015.	2.1	8
16	Prevalence and characteristics of a novel bovine adenovirus type 3 with a natural deletion fiber gene. <i>Infection, Genetics and Evolution</i> , 2020, 83, 104348.	2.3	7
17	Prevalence and genomic characteristics of a novel reassortment mammalian orthoreovirus type 2 in diarrhea piglets in Sichuan, China. <i>Infection, Genetics and Evolution</i> , 2020, 85, 104420.	2.3	7
18	<i>Staphylococcal enterotoxin M</i> induced inflammation and impairment of bovine mammary epithelial cells. <i>Journal of Dairy Science</i> , 2020, 103, 8350-8359.	3.4	30

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19	First detection and genomic characteristics of bovine torovirus in dairy calves in China. Archives of Virology, 2020, 165, 1577-1583.	2.1	7
20	High prevalence and genomic characteristics of G6P[1] Bovine Rotavirus A in yak in China. Journal of General Virology, 2020, 101, 701-711.	2.9	13
21	First detection of neboviruses in yak (<i>Bos grunniens</i>) and identification of a novel neboviruses based on complete genome. Veterinary Microbiology, 2019, 236, 108388.	1.9	6
22	Emergence of porcine epidemic diarrhea viruses with the novel S genes in Tibetan pigs in the Qinghai-Tibetan plateau in China. Virus Research, 2019, 270, 197652.	2.2	9
23	Molecular detection and genomic characteristics of bovine kobuvirus from dairy calves in China. Infection, Genetics and Evolution, 2019, 74, 103939.	2.3	13
24	Prevalence and complete genome of bovine norovirus with novel VP1 genotype in calves in China. Scientific Reports, 2019, 9, 12023.	3.3	6
25	Hepatitis E virus in Tibetan pigs on the Qinghai-Tibetan Plateau: detection, molecular characterization and infection of BALB/c mice. Archives of Virology, 2019, 164, 2943-2951.	2.1	6
26	A reverse transcription-insulated isothermal PCR assay for the detection of duck hepatitis A virus type 3 based on the POCKITâ„¢ system. Journal of Virological Methods, 2019, 270, 126-130.	2.1	3
27	Prevalence of a novel bovine coronavirus strain with a recombinant hemagglutinin/esterase gene in dairy calves in China. Transboundary and Emerging Diseases, 2019, 66, 1971-1981.	3.0	22
28	Surface-exposed loops L7 and L8 of Haemophilus (Glaesserella) parasuis OmpP2 contribute to the expression of proinflammatory cytokines in porcine alveolar macrophages. Veterinary Research, 2019, 50, 105.	3.0	7
29	A novel recombinant genome of minute virus of canines in China. Archives of Virology, 2019, 164, 861-865.	2.1	1
30	Detection and molecular characteristics of neboviruses in dairy cows in China. Journal of General Virology, 2019, 100, 35-45.	2.9	18
31	First detection of bovine coronavirus in Yak (<i>Bos grunniens</i>) and a bovine coronavirus genome with a recombinant HE gene. Journal of General Virology, 2019, 100, 793-803.	2.9	16
32	Genomic characterization of a RdRp-recombinant nebovirus strain with a novel VP1 genotype. Virus Research, 2018, 251, 6-13.	2.2	8
33	First detection of Nebovirus and Norovirus from cattle in China. Archives of Virology, 2018, 163, 475-478.	2.1	19
34	Prevalence and genomic characteristics of canine kobuvirus in southwest China. Archives of Virology, 2018, 163, 459-466.	2.1	14
35	Transcriptomic analysis reveals that enterovirus F strain SWUN-AB001 infection activates JNK/SAPK and p38 MAPK signaling pathways in MDBK cells. BMC Veterinary Research, 2018, 14, 395.	1.9	2
36	Prevalence and genome characteristics of canine astrovirus in southwest China. Journal of General Virology, 2018, 99, 880-889.	2.9	17

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37	IgtF effects of Haemophilus parasuis LOS induced inflammation through regulation of NF- κ B and MAPKs signaling pathways. Microbial Pathogenesis, 2017, 110, 380-384.	2.9	4
38	Isolation and characterization of a new enterovirus F in yak feces in the Qinghai-Tibetan Plateau. Archives of Virology, 2017, 162, 523-527.	2.1	10
39	A novel astrovirus species in the gut of yaks with diarrhoea in the Qinghai-Tibetan Plateau, 2013. Journal of General Virology, 2015, 96, 3672-3680.	2.9	26
40	Comparison of Culture Methods for Isolation of Salmonella in Yak Fecal Samples. Indian Journal of Microbiology, 2014, 54, 223-226.	2.7	4
41	Viral metagenomics analysis demonstrates the diversity of viral flora in piglet diarrhoeic faeces in China. Journal of General Virology, 2014, 95, 1603-1611.	2.9	98
42	Update on the pathogenesis of Haemophilus parasuis infection and virulence factors. Veterinary Microbiology, 2014, 168, 1-7.	1.9	47
43	Development of a Real-Time Quantitative PCR for Detecting Duck Hepatitis A Virus Genotype C. Journal of Clinical Microbiology, 2012, 50, 3318-3323.	3.9	22
44	Genetic and immunological characterization of G9 group porcine rotaviruses in China. Zoonoses and Public Health, 0, , .	2.2	1