

# Cheng Tang

## List of Publications by Year in descending order

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44  
papers

504  
citations

759233

12  
h-index

752698

20  
g-index

44  
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44  
docs citations

44  
times ranked

572  
citing authors

#	ARTICLE	IF	CITATIONS
1	Viral metagenomics analysis demonstrates the diversity of viral flora in piglet diarrhoeic faeces in China. <i>Journal of General Virology</i> , 2014, 95, 1603-1611.	2.9	98
2	Update on the pathogenesis of <i>Haemophilus parasuis</i> infection and virulence factors. <i>Veterinary Microbiology</i> , 2014, 168, 1-7.	1.9	47
3	Staphylococcal enterotoxin M induced inflammation and impairment of bovine mammary epithelial cells. <i>Journal of Dairy Science</i> , 2020, 103, 8350-8359.	3.4	30
4	A novel astrovirus species in the gut of yaks with diarrhoea in the Qinghai-Tibetan Plateau, 2013. <i>Journal of General Virology</i> , 2015, 96, 3672-3680.	2.9	26
5	Development of a Real-Time Quantitative PCR for Detecting Duck Hepatitis A Virus Genotype C. <i>Journal of Clinical Microbiology</i> , 2012, 50, 3318-3323.	3.9	22
6	Prevalence of a novel bovine coronavirus strain with a recombinant hemagglutinin/esterase gene in dairy calves in China. <i>Transboundary and Emerging Diseases</i> , 2019, 66, 1971-1981.	3.0	22
7	First detection of Nebovirus and Norovirus from cattle in China. <i>Archives of Virology</i> , 2018, 163, 475-478.	2.1	19
8	Detection and molecular characteristics of neboviruses in dairy cows in China. <i>Journal of General Virology</i> , 2019, 100, 35-45.	2.9	18
9	Prevalence and genome characteristics of canine astrovirus in southwest China. <i>Journal of General Virology</i> , 2018, 99, 880-889.	2.9	17
10	First detection of bovine coronavirus in Yak ( <i>Bos grunniens</i> ) and a bovine coronavirus genome with a recombinant HE gene. <i>Journal of General Virology</i> , 2019, 100, 793-803.	2.9	16
11	Prevalence and genomic characteristics of canine kobuvirus in southwest China. <i>Archives of Virology</i> , 2018, 163, 459-466.	2.1	14
12	Molecular detection and genomic characteristics of bovine kobuvirus from dairy calves in China. <i>Infection, Genetics and Evolution</i> , 2019, 74, 103939.	2.3	13
13	High prevalence and genomic characteristics of G6P[1] Bovine Rotavirus A in yak in China. <i>Journal of General Virology</i> , 2020, 101, 701-711.	2.9	13
14	Isolation and characterization of a new enterovirus F in yak feces in the Qinghai-Tibetan Plateau. <i>Archives of Virology</i> , 2017, 162, 523-527.	2.1	10
15	Emergence of porcine epidemic diarrhea viruses with the novel S genes in Tibetan pigs in the Qinghai-Tibetan plateau in China. <i>Virus Research</i> , 2019, 270, 197652.	2.2	9
16	Genomic characterization of a RdRp-recombinant nebovirus strain with a novel VP1 genotype. <i>Virus Research</i> , 2018, 251, 6-13.	2.2	8
17	First detection and molecular characteristics of caprine kobuvirus in goats in China. <i>Infection, Genetics and Evolution</i> , 2020, 85, 104566.	2.3	8
18	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

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19	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
20	Identification of a novel Aichivirus D in sheep. <i>Infection, Genetics and Evolution</i> , 2021, 91, 104810.	2.3	8
21	Surface-exposed loops L7 and L8 of <i>Haemophilus (Glaesserella) parasuis</i> OmpP2 contribute to the expression of proinflammatory cytokines in porcine alveolar macrophages. <i>Veterinary Research</i> , 2019, 50, 105.	3.0	7
22	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
23	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
24	First detection and genomic characteristics of bovine torovirus in dairy calves in China. <i>Archives of Virology</i> , 2020, 165, 1577-1583.	2.1	7
25	First detection of neboviruses in yak ( <i>Bos grunniens</i> ) and identification of a novel neboviruses based on complete genome. <i>Veterinary Microbiology</i> , 2019, 236, 108388.	1.9	6
26	Prevalence and complete genome of bovine norovirus with novel VP1 genotype in calves in China. <i>Scientific Reports</i> , 2019, 9, 12023.	3.3	6
27	Hepatitis E virus in Tibetan pigs on the Qinghai-Tibetan Plateau: detection, molecular characterization and infection of BALB/c mice. <i>Archives of Virology</i> , 2019, 164, 2943-2951.	2.1	6
28	Identification of a novel astrovirus in goats in China. <i>Infection, Genetics and Evolution</i> , 2021, 96, 105105.	2.3	5
29	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
30	Comparison of Culture Methods for Isolation of Salmonella in Yak Fecal Samples. <i>Indian Journal of Microbiology</i> , 2014, 54, 223-226.	2.7	4
31	IgtF effects of <i>Haemophilus parasuis</i> LOS induced inflammation through regulation of NF- $\kappa$ B and MAPKs signaling pathways. <i>Microbial Pathogenesis</i> , 2017, 110, 380-384.	2.9	4
32	Aichivirus C isolate is a diarrhoea-causing pathogen in goats. <i>Transboundary and Emerging Diseases</i> , 2022, 69, .	3.0	4
33	A reverse transcription-insulated isothermal PCR assay for the detection of duck hepatitis A virus type 3 based on the POCKIT $\hat{c}$ system. <i>Journal of Virological Methods</i> , 2019, 270, 126-130.	2.1	3
34	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
35	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
36	First detection of canine norovirus in dogs and a complete CVI.2 genome in mainland China. <i>Infection, Genetics and Evolution</i> , 2021, 92, 104879.	2.3	3

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37	First Detection and Genomic Characterization of Bovine Norovirus from Yak. <i>Pathogens</i> , 2022, 11, 192.	2.8	3
38	Transcriptomic analysis reveals that enterovirus F strain SWUN-AB001 infection activates JNK/SAPK and p38 MAPK signaling pathways in MDBK cells. <i>BMC Veterinary Research</i> , 2018, 14, 395.	1.9	2
39	Diversity of Astrovirus in Goats in Southwest China and Identification of Two Novel Caprine Astroviruses. <i>Microbiology Spectrum</i> , 2022, 10, .	3.0	2
40	A novel recombinant genome of minute virus of canines in China. <i>Archives of Virology</i> , 2019, 164, 861-865.	2.1	1
41	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
42	Genetic and immunological characterization of <scp>G9</scp> group AÂporcine rotaviruses in China. <i>Zoonoses and Public Health</i> , 0, , .	2.2	1
43	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
44	Identification and genomic characterization of bovine parvovirus 1 in yaks. <i>Journal of Veterinary Diagnostic Investigation</i> , 2021, 33, 104063872110296.	1.1	0