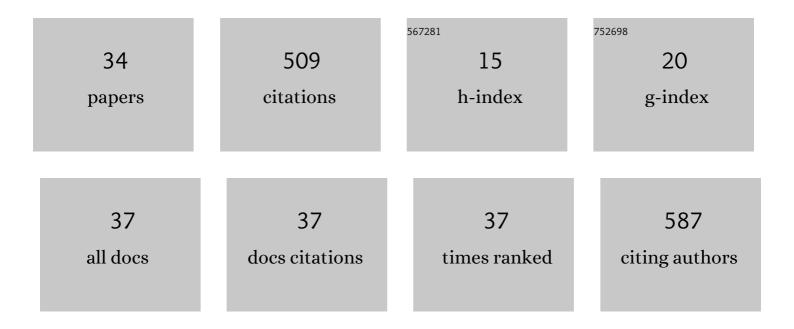
jinyan Gu

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Coronaviruses Nsp5 Antagonizes Porcine Gasdermin D-Mediated Pyroptosis by Cleaving Pore-Forming p30 Fragment. MBio, 2022, 13, e0273921.	4.1	28
2	TRAF6 autophagic degradation by <i>avibirnavirus</i> VP3 inhibits antiviral innate immunity via blocking NFKB/NF-κB activation. Autophagy, 2022, 18, 2781-2798.	9.1	18
3	PDPK1 regulates autophagosome biogenesis by binding to PIK3C3. Autophagy, 2021, 17, 2166-2183.	9.1	23
4	The serine-48 residue of nucleolar phosphoprotein nucleophosmin-1 plays critical role in subcellular localization and interaction with porcine circovirus type 3 capsid protein. Veterinary Research, 2021, 52, 4.	3.0	12
5	A Previously Undiscovered Circular RNA, circTNFAIP3, and Its Role in Coronavirus Replication. MBio, 2021, 12, e0298421.	4.1	10
6	Identification of functional IncRNAs in pseudorabies virus type II infected cells. Veterinary Microbiology, 2020, 242, 108564.	1.9	8
7	Nucleolar protein NPM1 is essential for circovirus replication by binding to viral capsid. Virulence, 2020, 11, 1379-1393.	4.4	16
8	Molecular characterization of an emerging reassortant mammalian orthoreovirus in China. Archives of Virology, 2020, 165, 2367-2372.	2.1	5
9	Conformational Dynamics of Nonenveloped Circovirus Capsid to the Host Cell Receptor. IScience, 2020, 23, 101547.	4.1	2
10	Porcine Epidemic Diarrhea Virus Deficient in RNA Cap Guanine-N-7 Methylation Is Attenuated and Induces Higher Type I and III Interferon Responses. Journal of Virology, 2020, 94, .	3.4	23
11	Protein Interactions Network of Porcine Circovirus Type 2 Capsid With Host Proteins. Frontiers in Microbiology, 2020, 11, 1129.	3.5	12
12	Antiviral Effect of Epigallocatechin Gallate via Impairing Porcine Circovirus Type 2 Attachment to Host Cell Receptor. Viruses, 2020, 12, 176.	3.3	21
13	NAP1L4 inhibits porcine circovirus type 2 replication via IFN-β signaling pathway. Veterinary Microbiology, 2020, 246, 108692.	1.9	7
14	Cross-species transmission resulted in the emergence and establishment of circovirus in pig. Infection, Genetics and Evolution, 2019, 75, 103973.	2.3	6
15	Conformational Changes and Nuclear Entry of Porcine Circovirus without Disassembly. Journal of Virology, 2019, 93, .	3.4	6
16	Genome Characteristics and Evolution of Pseudorabies Virus Strains in Eastern China from 2017 to 2019. Virologica Sinica, 2019, 34, 601-609.	3.0	26
17	Transcriptome Profiling Reveals Differential Effect of Interleukin-17A Upon Influenza Virus Infection in Human Cells. Frontiers in Microbiology, 2019, 10, 2344.	3.5	12
18	Detection and spike gene characterization in porcine deltacoronavirus in China during 2016–2018. Infection, Genetics and Evolution, 2019, 73, 151-158.	2.3	21

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#	Article	IF	CITATIONS
19	Characterization and epidemiological survey of porcine sapelovirus in China. Veterinary Microbiology, 2019, 232, 13-21.	1.9	14
20	Genetic diversity of porcine circovirus type 2 in China between 1999–2017. Transboundary and Emerging Diseases, 2019, 66, 599-605.	3.0	19
21	Comprehensive Genomic Characterization Analysis of IncRNAs in Cells With Porcine Delta Coronavirus Infection. Frontiers in Microbiology, 2019, 10, 3036.	3.5	7
22	Three amino acid substitutions in the NS1 protein change the virus replication of H5N1 influenza virus in human cells. Virology, 2018, 519, 64-73.	2.4	16
23	Molecular characteristics of the spike gene of porcine epidemic diarrhoea virus strains in Eastern China in 2016. Virus Research, 2018, 247, 47-54.	2.2	30
24	Caspase-Dependent Apoptosis Induction via Viral Protein ORF4 of Porcine Circovirus 2 Binding to Mitochondrial Adenine Nucleotide Translocase 3. Journal of Virology, 2018, 92, .	3.4	27
25	BECN1-dependent CASP2 incomplete autophagy induction by binding to rabies virus phosphoprotein. Autophagy, 2017, 13, 739-753.	9.1	45
26	Cellular proteomic analysis of porcine circovirus type 2 and classical swine fever virus coinfection in porcine kidneyâ€15 cells using isobaric tags for relative and absolute quantitationâ€coupled LCâ€MS/MS. Electrophoresis, 2017, 38, 1276-1291.	2.4	16
27	Identification and function analysis of canine stimulator of interferon gene (STING). Microbial Pathogenesis, 2017, 113, 202-208.	2.9	6
28	Characterization of H7N2 Avian Influenza Virus in Wild Birds and Pikas in Qinghai-Tibet Plateau Area. Scientific Reports, 2016, 6, 30974.	3.3	18
29	Human infections by avian influenza virus H5N6: Increasing risk by dynamic reassortment?. Infection, Genetics and Evolution, 2016, 42, 46-48.	2.3	2
30	Characterization of specific antigenic epitopes and the nuclear export signal of the Porcine circovirus 2 ORF3 protein. Veterinary Microbiology, 2016, 184, 40-50.	1.9	6
31	The A66G back mutation in NS2A of JEV SA 14 -14-2 strain contributes to production of NS1′ protein and the secreted NS1′ can be used for diagnostic biomarker for virulent virus infection. Infection, Genetics and Evolution, 2015, 36, 116-125.	2.3	12
32	In Vitro Coinfection and Replication of Classical Swine Fever Virus and Porcine Circovirus Type 2 in PK15 Cells. PLoS ONE, 2015, 10, e0139457.	2.5	12
33	Functional analysis of the interferon-stimulated response element of porcine circovirus type 2 and its role during viral replication in vitro and in vivo. Virology Journal, 2012, 9, 152.	3.4	13
34	Deletion of the single putative N-glycosylation site of the porcine circovirus type 2 Cap protein enhances specific immune responses by DNA immunisation in mice. Veterinary Journal, 2012, 192, 385-389.	1.7	8