Rui Wu

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1336887/publications.pdf

Version: 2024-02-01

		566801	713013
69	778	15	21
papers	citations	h-index	g-index
73	73	73	804
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Development and application of a visual microarray for synchronously detecting H5N1, H7N9 and H9N2 avian influenza virus RNA. Journal of Virological Methods, 2022, 301, 114371.	1.0	4
2	Molecular characterization of antimicrobial resistance and virulence factors of Enterococcus faecalis from ducks at slaughterhouses. Poultry Science, 2022, 101, 101646.	1.5	4
3	Study of the inhibitory effect of STAT1 on PDCoV infection. Veterinary Microbiology, 2022, 266, 109333.	0.8	7
4	HSP90 inhibitors 17-AAG and VER-82576 inhibit porcine deltacoronavirus replication in vitro. Veterinary Microbiology, 2022, 265, 109316.	0.8	6
5	Innate and mild Th17 cutaneous immune responses elicited by subcutaneous infection of immunocompetent mice with Cladosporium cladosporioides. Microbial Pathogenesis, 2022, 163, 105384.	1.3	2
6	Porcine Deltacoronavirus (PDCoV) Entry into PK-15 Cells by Caveolae-Mediated Endocytosis. Viruses, 2022, 14, 496.	1.5	6
7	A Comparative Transcriptomic Analysis Reveals That HSP90AB1 Is Involved in the Immune and Inflammatory Responses to Porcine Deltacoronavirus Infection. International Journal of Molecular Sciences, 2022, 23, 3280.	1.8	8
8	Phylogeny, Evolution, and Transmission Dynamics of Canine and Feline Coronaviruses: A Retro-Prospective Study. Frontiers in Microbiology, 2022, 13, 850516.	1.5	1
9	Deletion of Polyamine Transport Protein PotD Exacerbates Virulence in Glaesserella (Haemophilus) parasuis in the Form of Non-biofilm-generated Bacteria in a Murine Acute Infection Model. Virulence, 2021, 12, 520-546.	1.8	10
10	Identification and pathogenicity of <i>Plesiomonas shigelloides</i> from <i>Acipenser dabryanus</i> in China. Aquaculture Research, 2021, 52, 2286-2293.	0.9	12
11	Molecular and functional characterization of HtrA protein in Actinobacillus pleuropneumoniae. Veterinary Microbiology, 2021, 257, 109058.	0.8	4
12	Skin Microbiota of the Captive Giant Panda (Ailuropoda Melanoleuca) and the Distribution of Opportunistic Skin Disease-Associated Bacteria in Different Seasons. Frontiers in Veterinary Science, 2021, 8, 666486.	0.9	3
13	Regulatory effect of m ⁶ A modification on different viruses. Journal of Medical Virology, 2021, 93, 6100-6115.	2.5	7
14	Assessment of the pulmonary adaptive immune response to Cladosporium cladosporioides infection using an experimental mouse model. Scientific Reports, 2021, 11, 909.	1.6	8
15	Comparative transcriptome analysis reveals that deletion of CheY influences gene expressions of ABC transports and metabolism in Haemophilus parasuis. Functional and Integrative Genomics, 2021, 21, 695-707.	1.4	4
16	Skin Mycobiota of the Captive Giant Panda (Ailuropoda melanoleuca) and the Distribution of Opportunistic Dermatomycosis-Associated Fungi in Different Seasons. Frontiers in Veterinary Science, 2021, 8, 708077.	0.9	3
17	Identification of the immunodominant neutralizing regions in the spike glycoprotein of porcine deltacoronavirus. Virus Research, 2020, 276, 197834.	1.1	30
18	A class â lentogenic newcastle disease virus strain confers effective protection against the prevalent strains. Biologicals, 2020, 63, 74-80.	0.5	3

#	Article	IF	Citations
19	Evolutionary dynamics and transmission patterns of Newcastle disease virus in China through Bayesian phylogeographical analysis. PLoS ONE, 2020, 15, e0239809.	1.1	5
20	Identification of a Novel Linear B-Cell Epitope on the Nucleocapsid Protein of Porcine Deltacoronavirus. International Journal of Molecular Sciences, 2020, 21, 648.	1.8	18
21	Enhanced Immune Responses Against Japanese Encephalitis Virus Infection Using Japanese Encephalitis Live-Attenuated Virus Adjuvanted with Montanide GEL 01 ST in Mice. Vector-Borne and Zoonotic Diseases, 2019, 19, 835-843.	0.6	8
22	Comparison of Pathogenicity and Transmissibility of Influenza B and D Viruses in Pigs. Viruses, 2019, 11 , 905.	1.5	16
23	Aerosol and Contact Transmission Following Intranasal Infection of Mice with Japanese Encephalitis Virus. Viruses, 2019, 11, 87.	1.5	17
24	Acute oral toxicity test and assessment of combined toxicity of cadmium and aflatoxin B1 in kunming mice. Food and Chemical Toxicology, 2019, 131, 110577.	1.8	26
25	A requirement of TolC1 for effective survival, colonization and pathogenicity of Actinobacillus pleuropneumoniae. Microbial Pathogenesis, 2019, 134, 103596.	1.3	19
26	Identification, genotyping, and pathogenicity of Trichosporon spp. Isolated from Giant pandas (Ailuropoda melanoleuca). BMC Microbiology, 2019, 19, 113.	1.3	9
27	Construction of targeted and integrative promoter-reporter plasmids pDK-K and pDK-G to measure gene expression activity in Haemophilus parasuis. Microbial Pathogenesis, 2019, 134, 103565.	1.3	5
28	Hsp40 Protein DNAJB6 Interacts with Viral NS3 and Inhibits the Replication of the Japanese Encephalitis Virus. International Journal of Molecular Sciences, 2019, 20, 5719.	1.8	11
29	Characterization and Pathogenicity of the Porcine Deltacoronavirus Isolated in Southwest China. Viruses, 2019, 11, 1074.	1.5	32
30	Polyamine Transport Protein PotD Protects Mice against Haemophilus parasuis and Elevates the Secretion of Pro-Inflammatory Cytokines of Macrophage via JNK–MAPK and NF–κB Signal Pathways through TLR4. Vaccines, 2019, 7, 216.	2.1	7
31	Antigenic and Pathogenic Characteristics of QX-Type Avian Infectious Bronchitis Virus Strains Isolated in Southwestern China. Viruses, 2019, 11, 1154.	1.5	14
32	Mutation of I176R in the E coding region weakens Japanese encephalitis virus neurovirulence, but not its growth rate in BHK-21 cells. Archives of Virology, 2018, 163, 1351-1355.	0.9	13
33	Genomic changes in an attenuated genotype I Japanese encephalitis virus and comparison with virulent parental strain. Virus Genes, 2018, 54, 424-431.	0.7	3
34	OxyR of Haemophilus parasuis is a global transcriptional regulator important in oxidative stress resistance and growth. Gene, 2018, 643, 107-116.	1.0	12
35	Effective Pro-Inflammatory Induced Activity of GALT, a Conserved Antigen in A. Pleuropneumoniae, Improves the Cytokines Secretion of Macrophage via p38, ERK1/2 and JNK MAPKs Signal Pathway. Frontiers in Cellular and Infection Microbiology, 2018, 8, 337.	1.8	6
36	Serological and molecular epidemiology of Japanese encephalitis virus infections in swine herds in China, 2006–2012. Journal of Veterinary Science, 2018, 19, 151.	0.5	17

#	Article	IF	Citations
37	Basic Characterization of Natural Transformation in a Highly Transformable Haemophilus parasuis Strain SC1401. Frontiers in Cellular and Infection Microbiology, 2018, 8, 32.	1.8	10
38	QseC Mediates Osmotic Stress Resistance and Biofilm Formation in Haemophilus parasuis. Frontiers in Microbiology, 2018, 9, 212.	1.5	29
39	Galactose-1-phosphate uridyltransferase (GalT), an in vivo-induced antigen of Actinobacillus pleuropneumoniae serovar 5b strain L20, provided immunoprotection against serovar 1 strain MS71. PLoS ONE, 2018, 13, e0198207.	1.1	0
40	A trivalent Apx-fusion protein delivered by E. coli outer membrane vesicles induce protection against Actinobacillus pleuropneumoniae of serotype 1 and 7 challenge in a murine model. PLoS ONE, 2018, 13, e0191286.	1.1	8
41	Escherichia coli-derived outer membrane vesicles deliver galactose-1-phosphate uridyltransferase and yield partial protection against Actinobacillus pleuropneumoniae in mice. Journal of Microbiology and Biotechnology, 2018, 28, 2095-2105.	0.9	6
42	A streptomycin resistance marker in <i>H.Âparasuis</i> based on site-directed mutations in <i>rpsL</i> gene to perform unmarked in-frame mutations and to verify natural transformation. PeerJ, 2018, 6, e4253.	0.9	5
43	Introducing a cleavable signal peptide enhances the packaging efficiency of lentiviral vectors pseudotyped with Japanese encephalitis virus envelope proteins. Virus Research, 2017, 229, 9-16.	1.1	10
44	Polyamine-binding protein PotD2 is required for stress tolerance and virulence in Actinobacillus pleuropneumoniae. Antonie Van Leeuwenhoek, 2017, 110, 1647-1657.	0.7	6
45	TolC2 is required for the resistance, colonization and virulence of Actinobacillus pleuropneumoniae. Journal of Medical Microbiology, 2017, 66, 1170-1176.	0.7	10
46	Prevalence and seroepidemiology of <i>Haemophilus parasuis </i> in Sichuan province, China. PeerJ, 2017, 5, e3379.	0.9	18
47	A TolC-Like Protein of Actinobacillus pleuropneumoniae Is Involved in Antibiotic Resistance and Biofilm Formation. Frontiers in Microbiology, 2016, 07, 1618.	1.5	27
48	Immunoprotective Efficacy of Six In vivo-Induced Antigens against Actinobacillus pleuropneumoniae as Potential Vaccine Candidates in Murine Model. Frontiers in Microbiology, 2016, 7, 1623.	1.5	11
49	Absence of TolC Impairs Biofilm Formation in Actinobacillus pleuropneumoniae by Reducing Initial Attachment. PLoS ONE, 2016, 11, e0163364.	1.1	8
50	HtrA Is Important for Stress Resistance and Virulence in Haemophilus parasuis. Infection and Immunity, 2016, 84, 2209-2219.	1.0	35
51	Complete Genome Sequence of Highly Virulent Haemophilus parasuis Serotype 11 Strain SC1401. Genome Announcements, 2016, 4, .	0.8	9
52	Enhanced immune responses against Japanese encephalitis virus using recombinant adenoviruses coexpressing Japanese encephalitis virus envelope and porcine interleukin-6 proteins in mice. Virus Research, 2016, 222, 34-40.	1.1	6
53	Phylogenetic analysis reveals that Japanese encephalitis virus genotype III is still prevalent in swine herds in Sichuan province in China. Archives of Virology, 2016, 161, 1719-1722.	0.9	6
54	The NS3 and NS4A genes as the targets of RNA interference inhibit replication of Japanese encephalitis virus in vitro and in vivo. Gene, 2016, 594, 183-189.	1.0	8

#	Article	IF	CITATIONS
55	A recombinant nucleocapsid protein-based indirect enzyme-linked immunosorbent assay to detect antibodies against porcine deltacoronavirus. Journal of Veterinary Medical Science, 2016, 78, 601-606.	0.3	36
56	Immunogenicity of the recombinant HxuCBA proteins encoded by hxuCBA gene cluster of Haemophilus parasuis in mice. Gene, 2016, 591, 478-483.	1.0	7
57	The arcA gene contributes to the serum resistance and virulence of Haemophilus parasuis serovar 13 clinical strain EP3. Veterinary Microbiology, 2016, 196, 67-71.	0.8	20
58	Chryseobacterium chengduensis sp. nov. isolated from the air of captive giant panda enclosures in Chengdu, China. Journal of Zhejiang University: Science B, 2016, 17, 610-618.	1.3	4
59	Construction of a bivalent DNA vaccine co-expressing S genes of transmissible gastroenteritis virus and porcine epidemic diarrhea virus delivered by attenuated Salmonella typhimurium. Virus Genes, 2016, 52, 354-364.	0.7	24
60	Tissue tropism and molecular characterization of a Japanese encephalitis virus strain isolated from pigs in southwest China. Virus Research, 2016, 215, 55-64.	1.1	16
61	Immunogenicity of transmissible gastroenteritis virus (TGEV) M gene delivered by attenuated Salmonella typhimurium in mice. Virus Genes, 2016, 52, 218-227.	0.7	3
62	Effect of cheY deletion on growth and colonization in a Haemophilus parasuis serovar 13 clinical strain EP3. Gene, 2016, 577, 96-100.	1.0	22
63	Construction of an oral vaccine for transmissible gastroenteritis virus based on the TGEV N gene expressed in an attenuated Salmonella typhimurium vector. Journal of Virological Methods, 2016, 227, 6-13.	1.0	14
64	Promoter methylation, mRNA expression of goat tumor-associated genes and mRNA expression of DNA methyltransferase in enzootic nasal tumors. Molecular Medicine Reports, 2015, 12, 6275-6285.	1.1	0
65	Establishment of a Successive Markerless Mutation System in Haemophilus parasuis through Natural Transformation. PLoS ONE, 2015, 10, e0127393.	1.1	19
66	Identification of Actinobacillus pleuropneumoniae Genes Preferentially Expressed During Infection Using In Vivo-Induced Antigen Technology (IVIAT). Journal of Microbiology and Biotechnology, 2015, 25, 1606-1613.	0.9	5
67	Comparative proteomic analysis of the membrane proteins of two Haemophilus parasuis strains to identify proteins that may help in habitat adaptation and pathogenesis. Proteome Science, 2014, 12, 38.	0.7	9
68	First complete genomic characterization of a porcine parvovirus 5 isolate from China. Archives of Virology, 2014, 159, 1533-1536.	0.9	12
69	Comparative proteome analysis of the extracellular proteins of two Haemophilus parasuis strains Nagasaki and SW114. Biochemical and Biophysical Research Communications, 2014, 446, 997-1001.	1.0	12