## **Huigang Shen**

List of Publications by Year in descending order

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840776 713466 23 456 11 21 citations h-index g-index papers 23 23 23 770 times ranked docs citations citing authors all docs

#	Article	IF	CITATIONS
1	Domestic Pigs Are Susceptible to Infection with Influenza B Viruses. Journal of Virology, 2015, 89, 4818-4826.	3.4	73
2	High prevalence of porcine circovirus viremia in newborn piglets in five clinically normal swine breeding herds in North America. Preventive Veterinary Medicine, 2010, 97, 228-236.	1.9	55
3	Isolation of PCV3 from Perinatal and Reproductive Cases of PCV3-Associated Disease and In Vivo Characterization of PCV3 Replication in CD/CD Growing Pigs. Viruses, 2020, 12, 219.	3.3	46
4	Pathogenicity and transmissibility of reassortant H9 influenza viruses with genes from pandemic H1N1 virus. Journal of General Virology, 2012, 93, 2337-2345.	2.9	36
5	The neuraminidase and matrix genes of the 2009 pandemic influenza $H1N1$ virus cooperate functionally to facilitate efficient replication and transmissibility in pigs. Journal of General Virology, 2012, 93, 1261-1268.	2.9	36
6	Pathogenicity and Transmissibility of Novel Reassortant H3N2 Influenza Viruses with 2009 Pandemic H1N1 Genes in Pigs. Journal of Virology, 2015, 89, 2831-2841.	3.4	36
7	Evaluation of the efficacy of a commercial inactivated genogroup 2b-based porcine epidemic diarrhea virus (PEDV) vaccine and experimental live genogroup $1b$ exposure against $2b$ challenge. Veterinary Research, $2017$ , $48$ , $69$ .	3.0	28
8	Effect of porcine circovirus type 2a or 2b on infection kinetics and pathogenicity of two genetically divergent strains of porcine reproductive and respiratory syndrome virus in the conventional pig model. Veterinary Microbiology, 2012, 158, 69-81.	1.9	21
9	Association of concurrent porcine circovirus (PCV) 2a and 2b infection with PCV associated disease in vaccinated pigs. Research in Veterinary Science, 2013, 95, 775-781.	1.9	21
10	Role of enterotoxigenic <scp><i>Escherichia coli</i></scp> prophage in spreading antibiotic resistance in a porcineâ€derived environment. Environmental Microbiology, 2020, 22, 4974-4984.	3.8	21
11	Molecular cloning and characterization of Duck CD25. Veterinary Immunology and Immunopathology, 2007, 117, 266-274.	1.2	13
12	Porcine Hemagglutinating Encephalomyelitis Virus Infection <i>In Vivo</i> and <i>Ex Vivo</i> Journal of Virology, 2021, 95, .	3.4	13
13	Genetic diversity of porcine reproductive and respiratory syndrome virus $1$ in the United States of America from 2010 to 2018. Veterinary Microbiology, 2019, 239, 108486.	1.9	11
14	Detection and genomic characterization of new avian-like hepatitis E virus in a sparrow in the United States. Archives of Virology, 2018, 163, 2861-2864.	2.1	10
15	An interferon inducing porcine reproductive and respiratory syndrome virus vaccine candidate elicits protection against challenge with the heterologous virulent type 2 strain VR-2385 in pigs. Vaccine, 2017, 35, 125-131.	3.8	9
16	Virus survival and fitness when multiple genotypes and subtypes of influenza A viruses exist and circulate in swine. Virology, 2019, 532, 30-38.	2.4	8
17	In vivo CD4+ T-cell up-regulation and high dose side effects of refolded duck interleukin-2. Cytokine, 2006, 34, 297-302.	3.2	4
18	Genetic characterization of porcine sapoviruses identified from pigs during a diarrhoea outbreak in lowa, 2019. Transboundary and Emerging Diseases, 2022, 69, 1246-1255.	3.0	4

#	Article	IF	CITATIONS
19	In vitro and in vivo replication of influenza A H1N1 WSN33 viruses with different M1 proteins. Journal of General Virology, 2013, 94, 884-895.	2.9	3
20	Comparison of the efficacy of a commercial inactivated influenza A/H1N1/pdm09 virus (pH1N1) vaccine and two experimental M2e-based vaccines against pH1N1 challenge in the growing pig model. PLoS ONE, 2018, 13, e0191739.	2.5	3
21	Genomic Sequence of a Swine Pasivirus Type 1 Strain Identified in U.S. Swine. Genome Announcements, 2018, 6, .	0.8	2
22	Case Report and Genomic Characterization of a Novel Porcine Nodavirus in the United States. Viruses, 2021, 13, 73.	3.3	2
23	Genomic Sequence of a Megrivirus Strain Identified in Laying Hens in Brazil. Microbiology Resource Announcements, 2019, 8, .	0.6	1