## Sunhee Lee

## List of Publications by Year in descending order

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687363 794594 19 653 13 19 h-index citations g-index papers 19 19 19 654 citing authors docs citations times ranked all docs

#	Article	IF	CITATIONS
1	Complete Genome Characterization of Korean Porcine Deltacoronavirus Strain KOR/KNU14-04/2014. Genome Announcements, $2014, 2, .$	0.8	150
2	Outbreak-Related Porcine Epidemic Diarrhea Virus Strains Similar to US Strains, South Korea, 2013. Emerging Infectious Diseases, 2014, 20, 1223-1226.	4.3	138
3	Isolation and characterization of a Korean porcine epidemic diarrhea virus strain KNU-141112. Virus Research, 2015, 208, 215-224.	2.2	60
4	Robust and persistent SARS-CoV-2 infection in the human intestinal brush border expressing cells. Emerging Microbes and Infections, 2020, 9, 2169-2179.	6.5	43
5	Functional characterization and proteomic analysis of the nucleocapsid protein of porcine deltacoronavirus. Virus Research, 2015, 208, 136-145.	2.2	40
6	Genetic characteristics, pathogenicity, and immunogenicity associated with cell adaptation of a virulent genotype 2b porcine epidemic diarrhea virus. Veterinary Microbiology, 2017, 207, 248-258.	1.9	40
7	Efficacy of an inactivated genotype 2b porcine epidemic diarrhea virus vaccine in neonatal piglets. Veterinary Immunology and Immunopathology, 2016, 174, 45-49.	1.2	31
8	Full-Genome Sequence Analysis of a Variant Strain of Porcine Epidemic Diarrhea Virus in South Korea. Genome Announcements, $2014, 2, \ldots$	0.8	23
9	First detection of novel enterovirus G recombining a torovirus papainâ€like protease gene associated with diarrhoea in swine in South Korea. Transboundary and Emerging Diseases, 2019, 66, 1023-1028.	3.0	20
10	Reemergence of porcine epidemic diarrhea virus on Jeju Island. Korean Journal of Veterinary Research, 2014, 54, 185-188.	0.2	20
11	Genomic and antigenic characterization of porcine epidemic diarrhoea virus strains isolated from South Korea, 2017. Transboundary and Emerging Diseases, 2018, 65, 949-956.	3.0	18
12	Molecular characteristics and pathogenic assessment of porcine epidemic diarrhoea virus isolates from the 2018 endemic outbreaks on Jeju Island, South Korea. Transboundary and Emerging Diseases, 2019, 66, 1894-1909.	3.0	16
13	Complete genome sequence of a novel S-insertion variant of porcine epidemic diarrhea virus from South Korea. Archives of Virology, 2017, 162, 2919-2922.	2.1	14
14	Pathogenicity and genetic characteristics associated with cell adaptation of a virulent porcine reproductive and respiratory syndrome virus nsp2 DEL strain CA-2. Veterinary Microbiology, 2016, 186, 174-188.	1.9	9
15	Receptor-binding domain of SARS-CoV-2 spike protein efficiently inhibits SARS-CoV-2 infection and attachment to mouse lung. International Journal of Biological Sciences, 2021, 17, 3786-3794.	6.4	9
16	Complete genome sequence of a porcine astrovirus from South Korea. Archives of Virology, 2015, 160, 1819-1821.	2.1	8
17	A Novel Frameshifting Inhibitor Having Antiviral Activity against Zoonotic Coronaviruses. Viruses, 2021, 13, 1639.	3.3	7
18	Assessing the risk of recurrence of porcine epidemic diarrhea virus in affected farms on Jeju Island, South Korea. Journal of Veterinary Science, 2021, 22, e48.	1.3	6

#	ARTICLE	IF	CITATIONS
19	Phenotypic and genotypic analyses of an attenuated porcine reproductive and respiratory syndrome virus strain after serial passages in cultured porcine alveolar macrophages. Journal of Veterinary Science, 2018, 19, 358.	1.3	1