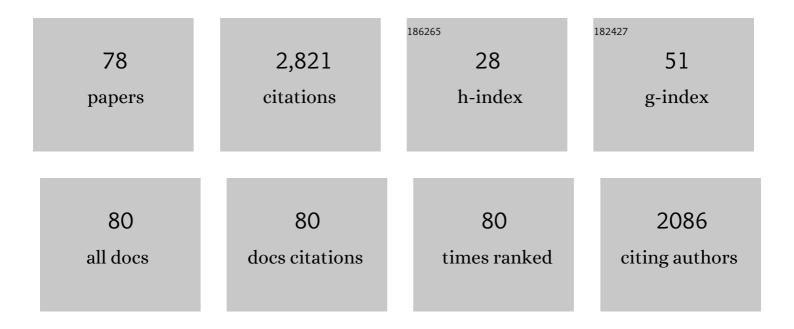
Changhee Lee

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Trypsin enhances SARS-CoV-2 infection by facilitating viral entry. Archives of Virology, 2022, 167, 441-458.	2.1	23
2	Complete genome sequence of a novel porcine hemagglutinating encephalomyelitis virus strain identified in South Korea. Archives of Virology, 2022, 167, 1381-1385.	2.1	2
3	Genetic and phylogenetic analysis of porcine circovirus type 2 on Jeju Island, South Korea, 2019–2020: evidence of a novel intergenotypic recombinant. Archives of Virology, 2021, 166, 1093-1102.	2.1	11
4	Inhibition of Antiviral Innate Immunity by Foot-and-Mouth Disease Virus L ^{pro} through Interaction with the N-Terminal Domain of Swine RNase L. Journal of Virology, 2021, 95, e0036121.	3.4	6
5	Successful Eradication of Porcine Epidemic Diarrhea in an Enzootically Infected Farm: A Two-Year Follow-Up Study. Pathogens, 2021, 10, 830.	2.8	14
6	Stress-activated protein kinases are involved in the replication of porcine deltacoronavirus. Virology, 2021, 559, 196-209.	2.4	5
7	A simple colorimetric detection of porcine epidemic diarrhea virus by reverse transcription loop-mediated isothermal amplification assay using hydroxynaphthol blue metal indicator. Journal of Virological Methods, 2021, 298, 114289.	2.1	12
8	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
9	Emergence and evolution of novel G2b-like porcine epidemic diarrhea virus inter-subgroup G1b recombinants. Archives of Virology, 2020, 165, 2471-2478.	2.1	11
10	Novel lineage 1 recombinants of porcine reproductive and respiratory syndrome virus isolated from vaccinated herds: genome sequences and cytokine production profiles. Archives of Virology, 2020, 165, 2259-2277.	2.1	10
11	Genomic characterization of classical swine fever virus LOM variants with 3′-UTR INDELs from pigs on Jeju Island, South Korea. Archives of Virology, 2020, 165, 1691-1696.	2.1	7
12	Advanced targetâ€specific probeâ€based realâ€time loopâ€mediated isothermal amplification assay for the rapid and specific detection of porcine circovirus 3. Transboundary and Emerging Diseases, 2020, 67, 2336-2344.	3.0	10
13	Probeâ€based realâ€time reverse transcription loopâ€mediated isothermal amplification (RRTâ€LAMP) assay for rapid and specific detection of footâ€andâ€mouth disease virus. Transboundary and Emerging Diseases, 2020, 67, 2936-2945.	3.0	8
14	Porcine deltacoronavirus activates the Raf/MEK/ERK pathway to promote its replication. Virus Research, 2020, 283, 197961.	2.2	12
15	Complete genome sequences of novel S-deletion variants of porcine epidemic diarrhea virus identified from a recurrent outbreak on Jeju Island, South Korea. Archives of Virology, 2019, 164, 2621-2625.	2.1	6
16	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
17	Assessment of the safety and efficacy of an attenuated live vaccine based on highly virulent genotype 2b porcine epidemic diarrhea virus in nursing piglets. Veterinary Microbiology, 2019, 231, 120-128.	1.9	19
18	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

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19	Endemic outbreaks due to the reâ€emergence of classical swine fever after accidental introduction of modified live LOM vaccine on Jeju Island, South Korea. Transboundary and Emerging Diseases, 2019, 66, 634-639.	3.0	12
20	Porcine Epidemic Diarrhea Virus. , 2019, , .		1
21	Functional Characterization and Proteomic Analysis of Porcine Deltacoronavirus Accessory Protein NS7. Journal of Microbiology and Biotechnology, 2019, 29, 1817-1829.	2.1	12
22	Generation and protective efficacy of a cold-adapted attenuated genotype 2b porcine epidemic diarrhea virus. Journal of Veterinary Science, 2019, 20, e32.	1.3	7
23	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
24	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
25	Porcine deltacoronavirus induces caspase-dependent apoptosis through activation of the cytochrome c-mediated intrinsic mitochondrial pathway. Virus Research, 2018, 253, 112-123.	2.2	57
26	Cholesterol is important for the entry process of porcine deltacoronavirus. Archives of Virology, 2018, 163, 3119-3124.	2.1	42
27	Geographic distribution and molecular analysis of porcine reproductive and respiratory syndrome viruses circulating in swine farms in the Republic of Korea between 2013 and 2016. BMC Veterinary Research, 2018, 14, 160.	1.9	31
28	Isolation and characterization of Korean porcine deltacoronavirus strain KNU16-07. Journal of Veterinary Science, 2018, 19, 577.	1.3	26
29	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
30	Cellular cholesterol is required for porcine nidovirus infection. Archives of Virology, 2017, 162, 3753-3767.	2.1	28
31	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
32	Prevalence, complete genome sequencing and phylogenetic analysis of porcine deltacoronavirus in South Korea, 2014-2016. Transboundary and Emerging Diseases, 2017, 64, 1364-1370.	3.0	59
33	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
34	JNK and p38 mitogen-activated protein kinase pathways contribute to porcine epidemic diarrhea virus infection. Virus Research, 2016, 222, 1-12.	2.2	38
35	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
36	The complete genome sequence of a lactic acid bacterium Leuconostoc mesenteroides ssp. dextranicum strain DSM 20484T. Journal of Biotechnology, 2016, 219, 3-4.	3.8	2

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37	lvermectin inhibits porcine reproductive and respiratory syndrome virus in cultured porcine alveolar macrophages. Archives of Virology, 2016, 161, 257-268.	2.1	19
38	Draft Genome Sequence of Caprolactam-Degrading Pseudomonas putida Strain SJ3. Genome Announcements, 2015, 3, .	0.8	2
39	Porcine epidemic diarrhea virus: An emerging and re-emerging epizootic swine virus. Virology Journal, 2015, 12, 193.	3.4	392
40	Sasa quelpaertensis Nakai extract suppresses porcine reproductive and respiratory syndrome virus replication and modulates virus-induced cytokine production. Archives of Virology, 2015, 160, 1977-1988.	2.1	9
41	Functional characterization and proteomic analysis of the nucleocapsid protein of porcine deltacoronavirus. Virus Research, 2015, 208, 136-145.	2.2	40
42	Isolation and characterization of a Korean porcine epidemic diarrhea virus strain KNU-141112. Virus Research, 2015, 208, 215-224.	2.2	60
43	Immunoprophylactic effect of chicken egg yolk antibody (IgY) against a recombinant S1 domain of the porcine epidemic diarrhea virus spike protein in piglets. Archives of Virology, 2015, 160, 2197-2207.	2.1	17
44	Extracellular signal-regulated kinase (ERK) activation is required for porcine epidemic diarrhea virus replication. Virology, 2015, 484, 181-193.	2.4	38
45	Complete genome sequence of a porcine astrovirus from South Korea. Archives of Virology, 2015, 160, 1819-1821.	2.1	8
46	Complete genome sequence of a keratin-degrading bacterium Chryseobacterium gallinarum strain DSM 27622T isolated from chicken. Journal of Biotechnology, 2015, 211, 66-67.	3.8	10
47	Outbreak-Related Porcine Epidemic Diarrhea Virus Strains Similar to US Strains, South Korea, 2013. Emerging Infectious Diseases, 2014, 20, 1223-1226.	4.3	138
48	Full-Genome Sequence Analysis of a Variant Strain of Porcine Epidemic Diarrhea Virus in South Korea. Genome Announcements, 2014, 2, .	0.8	23
49	Immunogenicity and protective efficacy of recombinant S1 domain of the porcine epidemic diarrhea virus spike protein. Archives of Virology, 2014, 159, 2977-2987.	2.1	89
50	Complete Genome Characterization of Korean Porcine Deltacoronavirus Strain KOR/KNU14-04/2014. Genome Announcements, 2014, 2, .	0.8	150
51	Genomic analysis and pathogenic characteristics of Type 2 porcine reproductive and respiratory syndrome virus nsp2 deletion strains isolated in Korea. Veterinary Microbiology, 2014, 170, 232-245.	1.9	19
52	Porcine epidemic diarrhea virus induces caspase-independent apoptosis through activation of mitochondrial apoptosis-inducing factor. Virology, 2014, 460-461, 180-193.	2.4	78
53	Reemergence of porcine epidemic diarrhea virus on Jeju Island. Korean Journal of Veterinary Research, 2014, 54, 185-188.	0.2	20
54	Complete genome sequence of a novel porcine parainfluenza virus 5 isolate in Korea. Archives of Virology, 2013, 158, 1765-1772.	2.1	49

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55	Characterization in vitro and in vivo of a novel porcine parainfluenza virus 5 isolate in Korea. Virus Research, 2013, 178, 423-429.	2.2	9
56	Ribavirin efficiently suppresses porcine nidovirus replication. Virus Research, 2013, 171, 44-53.	2.2	42
57	Cytokine production in immortalized porcine alveolar macrophages infected with porcine reproductive and respiratory syndrome virus. Veterinary Immunology and Immunopathology, 2012, 150, 213-220.	1.2	21
58	Proteomic characterization of a novel structural protein ORF5a of porcine reproductive and respiratory syndrome virus. Virus Research, 2012, 169, 255-263.	2.2	15
59	Human telomerase reverse transcriptase-immortalized porcine monomyeloid cell lines for the production of porcine reproductive and respiratory syndrome virus. Journal of Virological Methods, 2012, 179, 26-32.	2.1	22
60	Stress-activated protein kinases are involved in porcine reproductive and respiratory syndrome virus infection and modulate virus-induced cytokine production. Virology, 2012, 427, 80-89.	2.4	24
61	Porcine reproductive and respiratory syndrome virus nucleocapsid protein modulates interferon-β production by inhibiting IRF3 activation in immortalized porcine alveolar macrophages. Archives of Virology, 2011, 156, 2187-2195.	2.1	88
62	Deletion of the cytoplasmic domain of CD163 enhances porcine reproductive and respiratory syndrome virus replication. Archives of Virology, 2010, 155, 1319-1323.	2.1	13
63	Generation of a porcine alveolar macrophage cell line for the growth of porcine reproductive and respiratory syndrome virus. Journal of Virological Methods, 2010, 163, 410-415.	2.1	81
64	A molecular analysis of European porcine reproductive and respiratory syndrome virus isolated in South Korea. Veterinary Microbiology, 2010, 143, 394-400.	1.9	53
65	Contribution of the porcine aminopeptidase N (CD13) receptor density to porcine epidemic diarrhea virus infection. Veterinary Microbiology, 2010, 144, 41-50.	1.9	86
66	Development and characterization of stable cell lines constitutively expressing the porcine reproductive and respiratory syndrome virus nucleocapsid protein. Journal of Veterinary Science, 2010, 11, 169.	1.3	3
67	Heterogeneity in spike protein genes of porcine epidemic diarrhea viruses isolated in Korea. Virus Research, 2010, 149, 175-182.	2.2	111
68	Differential cellular protein expression in continuous porcine alveolar macrophages regulated by the porcine reproductive and respiratory syndrome virus nucleocapsid protein. Virus Research, 2010, 151, 88-96.	2.2	7
69	Porcine reproductive and respiratory syndrome virus replication is suppressed by inhibition of the extracellular signal-regulated kinase (ERK) signaling pathway. Virus Research, 2010, 152, 50-58.	2.2	44
70	Complete genomic characterization of a European type 1 porcine reproductive and respiratory syndrome virus isolate in Korea. Archives of Virology, 2009, 154, 629-638.	2.1	49
71	Genetic differentiation of the nucleocapsid protein of Korean isolates of porcine epidemic diarrhoea virus by RT-PCR based restriction fragment length polymorphism analysis. Veterinary Journal, 2008, 178, 138-140.	1.7	7
72	Mutations within the nuclear localization signal of the porcine reproductive and respiratory syndrome virus nucleocapsid protein attenuate virus replication. Virology, 2006, 346, 238-250.	2.4	82

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73	The small envelope protein of porcine reproductive and respiratory syndrome virus possesses ion channel protein-like properties. Virology, 2006, 355, 30-43.	2.4	73
74	The Nuclear Localization Signal of the Prrs Virus Nucleocapsid Protein Modulates Viral Replication in vitro and Antibody Response in vivo. Advances in Experimental Medicine and Biology, 2006, 581, 145-148.	1.6	12
75	A DNA-launched reverse genetics system for porcine reproductive and respiratory syndrome virus reveals that homodimerization of the nucleocapsid protein is essential for virus infectivity. Virology, 2005, 331, 47-62.	2.4	74
76	Cysteine residues of the porcine reproductive and respiratory syndrome virus small envelope protein are non-essential for virus infectivity. Journal of General Virology, 2005, 86, 3091-3096.	2.9	29
77	Infectious cDNA clones of porcine reproductive and respiratory syndrome virus and their potential as vaccine vectors. Veterinary Immunology and Immunopathology, 2004, 102, 143-154.	1.2	43
78	Differential host cell gene expression regulated by the porcine reproductive and respiratory syndrome virus GP4 and GP5 glycoproteins. Veterinary Immunology and Immunopathology, 2004, 102, 189-198.	1.2	25