

# Min-Suk Song

## List of Publications by Year in descending order

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72  
papers

3,231  
citations

236925

25  
h-index

168389

53  
g-index

77  
all docs

77  
docs citations

77  
times ranked

6096  
citing authors

#	ARTICLE	IF	CITATIONS
1	Infection and Rapid Transmission of SARS-CoV-2 in Ferrets. <i>Cell Host and Microbe</i> , 2020, 27, 704-709.e2.	11.0	815
2	Development of a reverse transcription-loop-mediated isothermal amplification as a rapid early-detection method for novel SARS-CoV-2. <i>Emerging Microbes and Infections</i> , 2020, 9, 998-1007.	6.5	267
3	Antiviral Efficacies of FDA-Approved Drugs against SARS-CoV-2 Infection in Ferrets. <i>MBio</i> , 2020, 11, .	4.1	165
4	The Polymerase Acidic Protein Gene of Influenza A Virus Contributes to Pathogenicity in a Mouse Model. <i>Journal of Virology</i> , 2009, 83, 12325-12335.	3.4	149
5	Rapid and simple colorimetric detection of multiple influenza viruses infecting humans using a reverse transcriptional loop-mediated isothermal amplification (RT-LAMP) diagnostic platform. <i>BMC Infectious Diseases</i> , 2019, 19, 676.	2.9	144
6	Continuing evolution of H9 influenza viruses in Korean poultry. <i>Virology</i> , 2007, 359, 313-323.	2.4	106
7	One-Pot Reverse Transcriptional Loop-Mediated Isothermal Amplification (RT-LAMP) for Detecting MERS-CoV. <i>Frontiers in Microbiology</i> , 2016, 7, 2166.	3.5	99
8	Mammalian adaptation of influenza A(H7N9) virus is limited by a narrow genetic bottleneck. <i>Nature Communications</i> , 2015, 6, 6553.	12.8	90
9	Evidence of Human-to-Swine Transmission of the Pandemic (H1N1) 2009 Influenza Virus in South Korea. <i>Journal of Clinical Microbiology</i> , 2010, 48, 3204-3211.	3.9	76
10	Ferret animal model of severe fever with thrombocytopenia syndrome phlebovirus for human lethal infection and pathogenesis. <i>Nature Microbiology</i> , 2019, 4, 438-446.	13.3	66
11	Sublingual Immunization with M2-Based Vaccine Induces Broad Protective Immunity against Influenza. <i>PLoS ONE</i> , 2011, 6, e27953.	2.5	66
12	Profiling and Characterization of Influenza Virus N1 Strains Potentially Resistant to Multiple Neuraminidase Inhibitors. <i>Journal of Virology</i> , 2015, 89, 287-299.	3.4	54
13	Critical role of neutralizing antibody for SARS-CoV-2 reinfection and transmission. <i>Emerging Microbes and Infections</i> , 2021, 10, 152-160.	6.5	54
14	Virulence and Genetic Compatibility of Polymerase Reassortant Viruses Derived from the Pandemic (H1N1) 2009 Influenza Virus and Circulating Influenza A Viruses. <i>Journal of Virology</i> , 2011, 85, 6275-6286.	3.4	51
15	Identification and characterization of influenza variants resistant to a viral endonuclease inhibitor. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 3669-3674.	7.1	51
16	Development of a SFTSV DNA vaccine that confers complete protection against lethal infection in ferrets. <i>Nature Communications</i> , 2019, 10, 3836.	12.8	51
17	Molecular characterization of avian influenza H5N1 virus in Egypt and the emergence of a novel endemic subclade. <i>Journal of General Virology</i> , 2014, 95, 1444-1463.	2.9	46
18	The Emergence and Decenary Distribution of Clade 2.3.4.4 HPAI H5Nx. <i>Microorganisms</i> , 2019, 7, 156.	3.6	46

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19	Genetic characterisation of novel, highly pathogenic avian influenza (HPAI) H5N6 viruses isolated in birds, South Korea, November 2016. <i>Eurosurveillance</i> , 2017, 22, .	7.0	44
20	Unique Determinants of Neuraminidase Inhibitor Resistance among N3, N7, and N9 Avian Influenza Viruses. <i>Journal of Virology</i> , 2015, 89, 10891-10900.	3.4	43
21	Ecology of H3 avian influenza viruses in Korea and assessment of their pathogenic potentials. <i>Journal of General Virology</i> , 2008, 89, 949-957.	2.9	42
22	Screening for Neuraminidase Inhibitor Resistance Markers among Avian Influenza Viruses of the N4, N5, N6, and N8 Neuraminidase Subtypes. <i>Journal of Virology</i> , 2018, 92, .	3.4	42
23	Mucosal immunity induced by adenovirus-based H5N1 HPAI vaccine confers protection against a lethal H5N2 avian influenza virus challenge. <i>Virology</i> , 2009, 395, 182-189.	2.4	37
24	Comparison of the pathogenic potential of highly pathogenic avian influenza (HPAI) H5N6, and H5N8 viruses isolated in South Korea during the 2016-2017 winter season. <i>Emerging Microbes and Infections</i> , 2018, 7, 1-10.	6.5	32
25	Age-dependent pathogenic characteristics of SARS-CoV-2 infection in ferrets. <i>Nature Communications</i> , 2022, 13, 21.	12.8	31
26	Molecular Markers for Interspecies Transmission of Avian Influenza Viruses in Mammalian Hosts. <i>International Journal of Molecular Sciences</i> , 2017, 18, 2706.	4.1	29
27	Molecular Signatures of Inflammatory Profile and B-Cell Function in Patients with Severe Fever with Thrombocytopenia Syndrome. <i>MBio</i> , 2021, 12, .	4.1	25
28	Simple, Rapid and Sensitive Portable Molecular Diagnosis of SFTS Virus Using Reverse Transcriptional Loop-Mediated Isothermal Amplification (RT-LAMP). <i>Journal of Microbiology and Biotechnology</i> , 2018, 28, 1928-1936.	2.1	25
29	Complete protection against a H5N2 avian influenza virus by a DNA vaccine expressing a fusion protein of H1N1 HA and M2e. <i>Vaccine</i> , 2011, 29, 5481-5487.	3.8	24
30	Bacterial Outer Membrane Vesicles Provide Broad-Spectrum Protection against Influenza Virus Infection via Recruitment and Activation of Macrophages. <i>Journal of Innate Immunity</i> , 2019, 11, 316-329.	3.8	24
31	Genetic and phylogenetic characterizations of a novel genotype of highly pathogenic avian influenza (HPAI) H5N8 viruses in 2016/2017 in South Korea. <i>Infection, Genetics and Evolution</i> , 2017, 53, 56-67.	2.3	23
32	Prokaryote-expressed M2e protein improves H9N2 influenza vaccine efficacy and protection against lethal influenza a virus in mice. <i>Virology Journal</i> , 2013, 10, 104.	3.4	21
33	Correlation Between the Interval of Influenza Virus Infectivity and Results of Diagnostic Assays in a Ferret Model. <i>Journal of Infectious Diseases</i> , 2016, 213, 407-410.	4.0	21
34	Investigation of the biological indicator for vaccine efficacy against highly pathogenic avian influenza (HPAI) H5N1 virus challenge in mice and ferrets. <i>Vaccine</i> , 2009, 27, 3145-3152.	3.8	20
35	Establishment of Vero cell RNA polymerase I-driven reverse genetics for Influenza A virus and its application for pandemic (H1N1) 2009 influenza virus vaccine production. <i>Journal of General Virology</i> , 2013, 94, 1230-1235.	2.9	20
36	Mouse adaptation of influenza B virus increases replication in the upper respiratory tract and results in droplet transmissibility in ferrets. <i>Scientific Reports</i> , 2015, 5, 15940.	3.3	20

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37	Vaccine Efficacy of Inactivated, Chimeric Hemagglutinin H9/H5N2 Avian Influenza Virus and Its Suitability for the Marker Vaccine Strategy. <i>Journal of Virology</i> , 2017, 91, .	3.4	18
38	Development of a dual-protective live attenuated vaccine against H5N1 and H9N2 avian influenza viruses by modifying the NS1 gene. <i>Archives of Virology</i> , 2015, 160, 1729-1740.	2.1	16
39	Cross-protective efficacies of highly-pathogenic avian influenza H5N1 vaccines against a recent H5N8 virus. <i>Virology</i> , 2016, 498, 36-43.	2.4	16
40	Molecular characterization of mammalian-adapted Korean-type avian H9N2 virus and evaluation of its virulence in mice. <i>Journal of Microbiology</i> , 2015, 53, 570-577.	2.8	15
41	Characterization of an H4N2 influenza virus from Quails with a multibasic motif in the hemagglutinin cleavage site. <i>Virology</i> , 2014, 468-470, 72-80.	2.4	14
42	Comparison of the virulence and transmissibility of canine H3N2 influenza viruses and characterization of their canine adaptation factors. <i>Emerging Microbes and Infections</i> , 2018, 7, 1-14.	6.5	14
43	Rapid acquisition of polymorphic virulence markers during adaptation of highly pathogenic avian influenza H5N8 virus in the mouse. <i>Scientific Reports</i> , 2017, 7, 40667.	3.3	13
44	Altered virulence of Highly Pathogenic Avian Influenza (HPAI) H5N8 reassortant viruses in mammalian models. <i>Virulence</i> , 2018, 9, 133-148.	4.4	13
45	Early Regulation of Viral Infection Reduces Inflammation and Rescues Mx-Positive Mice from Lethal Avian Influenza Infection. <i>American Journal of Pathology</i> , 2013, 182, 1308-1321.	3.8	11
46	Evaluation of the zoonotic potential of a novel reassortant H1N2 swine influenza virus with gene constellation derived from multiple viral sources. <i>Infection, Genetics and Evolution</i> , 2015, 34, 378-393.	2.3	11
47	Genetic characteristics of highly pathogenic H5N8 avian influenza viruses isolated from migratory wild birds in South Korea during 2014-2015. <i>Archives of Virology</i> , 2016, 161, 2749-2764.	2.1	11
48	An I436N substitution confers resistance of influenza A(H1N1)pdm09 viruses to multiple neuraminidase inhibitors without affecting viral fitness. <i>Journal of General Virology</i> , 2018, 99, 292-302.	2.9	11
49	Genetic diversity and pathogenic potential of low pathogenic H7 avian influenza viruses isolated from wild migratory birds in Korea. <i>Infection, Genetics and Evolution</i> , 2016, 45, 268-284.	2.3	10
50	Evaluation of the Immune Responses to and Cross-Protective Efficacy of Eurasian H7 Avian Influenza Viruses. <i>Journal of Virology</i> , 2017, 91, .	3.4	10
51	A Novel Neuraminidase-Dependent Hemagglutinin Cleavage Mechanism Enables the Systemic Spread of an H7N6 Avian Influenza Virus. <i>MBio</i> , 2019, 10, .	4.1	10
52	Baloxavir-oseltamivir combination therapy inhibits the emergence of resistant substitutions in influenza A virus PA gene in a mouse model. <i>Antiviral Research</i> , 2021, 193, 105126.	4.1	10
53	Development of a Multiplex RT-qPCR for the Detection of Different Clades of Avian Influenza in Poultry. <i>Viruses</i> , 2020, 12, 100.	3.3	9
54	The Homologous Tripartite Viral RNA Polymerase of A/Swine/Korea/CT1204/2009(H1N2) Influenza Virus Synergistically Drives Efficient Replication and Promotes Respiratory Droplet Transmission in Ferrets. <i>Journal of Virology</i> , 2013, 87, 10552-10562.	3.4	8

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55	Zika virus lateral flow assays using reverse transcription-loop-mediated isothermal amplification. <i>RSC Advances</i> , 2021, 11, 17800-17808.	3.6	8
56	Preclinical evaluation of the efficacy of an H5N8 vaccine candidate (IDCDC-RG43A) in mouse and ferret models for pandemic preparedness. <i>Vaccine</i> , 2019, 37, 484-493.	3.8	7
57	Virulence of pandemic (H1N1) 2009 influenza A polymerase reassortant viruses. <i>Virulence</i> , 2011, 2, 422-426.	4.4	6
58	Virological and pathological characterization of an avian H1N1 influenza A virus. <i>Archives of Virology</i> , 2018, 163, 1153-1162.	2.1	6
59	<i>In Vitro</i> and <i>In Vivo</i> Characterization of Novel Neuraminidase Substitutions in Influenza A(H1N1)pdm09 Virus Identified Using Laninamivir-Mediated <i>In Vitro</i> Selection. <i>Journal of Virology</i> , 2019, 93, .	3.4	6
60	Peptide Nucleic Acid (PNA)-Enhanced Specificity of a Dual-Target Real-Time Quantitative Polymerase Chain Reaction (RT-qPCR) Assay for the Detection and Differentiation of SARS-CoV-2 from Related Viruses. <i>Diagnostics</i> , 2020, 10, 775.	2.6	6
61	JEV-nanobarcode and colorimetric reverse transcription loop-mediated isothermal amplification (cRT-LAMP). <i>Mikrochimica Acta</i> , 2021, 188, 333.	5.0	6
62	Eyedrop Vaccination Induced Systemic and Mucosal Immunity against Influenza Virus in Ferrets. <i>PLoS ONE</i> , 2016, 11, e0157634.	2.5	5
63	The significance of avian influenza virus mouse-adaptation and its application in characterizing the efficacy of new vaccines and therapeutic agents. <i>Clinical and Experimental Vaccine Research</i> , 2017, 6, 83.	2.2	5
64	Evaluation of the efficacy of a pre-pandemic H5N1 vaccine (MG1109) in mouse and ferret models. <i>Journal of Microbiology</i> , 2012, 50, 478-488.	2.8	4
65	Growth and Pathogenic Potential of Naturally Selected Reassortants after Coinfection with Pandemic H1N1 and Highly Pathogenic Avian Influenza H5N1 Viruses. <i>Journal of Virology</i> , 2016, 90, 616-623.	3.4	4
66	Seroprevalence of Severe Fever with Thrombocytopenia Syndrome Phlebovirus in Domesticated Deer in South Korea. <i>Virologica Sinica</i> , 2019, 34, 501-507.	3.0	4
67	Development of a rapid, simple and efficient one-pot cloning method for a reverse genetics system of broad subtypes of influenza A virus. <i>Scientific Reports</i> , 2019, 9, 8318.	3.3	4
68	Development of a Rapid Fluorescent Diagnostic System for Early Detection of the Highly Pathogenic Avian Influenza H5 Clade 2.3.4.4 Viruses in Chicken Stool. <i>International Journal of Molecular Sciences</i> , 2022, 23, 6301.	4.1	4
69	Efficacy of A/H1N1/2009 split inactivated influenza A vaccine (GC1115) in mice and ferrets. <i>Journal of Microbiology</i> , 2019, 57, 163-169.	2.8	3
70	<i>In Vitro</i> Profiling of Laninamivir-Resistant Substitutions in N3 to N9 Avian Influenza Virus Neuraminidase Subtypes and Their Association with <i>In Vivo</i> Susceptibility. <i>Journal of Virology</i> , 2020, 95, .	3.4	3
71	Genetic Characteristics and Phylogenetic Analysis of Influenza Type B Viruses Isolated from Nasopharyngeal Suction Samples of Korean Patients. <i>Journal of Bacteriology and Virology</i> , 2009, 39, 125.	0.1	2
72	Multiple HA substitutions in highly pathogenic avian influenza H5Nx viruses contributed to the change in the NA subtype preference. <i>Virulence</i> , 2022, 13, 990-1004.	4.4	1