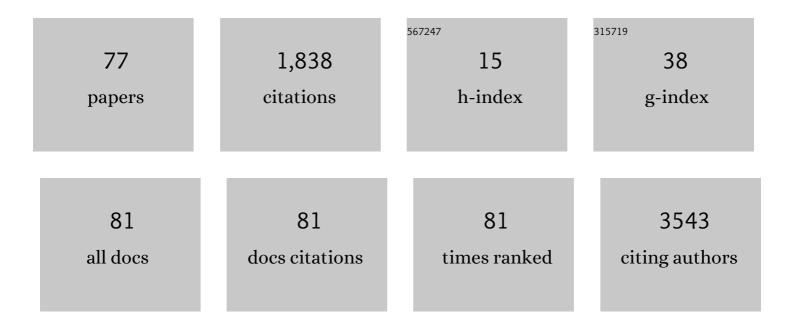
Nanping Wu

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/57074/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Molecular Architecture of the SARS-CoV-2 Virus. Cell, 2020, 183, 730-738.e13.	28.9	793
2	Alterations in the Fecal Microbiota of Patients with HIV-1 Infection: An Observational Study in A Chinese Population. Scientific Reports, 2016, 6, 30673.	3.3	153
3	Virus strain from a mild COVID-19 patient in Hangzhou represents a new trend in SARS-CoV-2 evolution potentially related to Furin cleavage site. Emerging Microbes and Infections, 2020, 9, 1474-1488.	6.5	51
4	Novel reassortant highly pathogenic H5N6 avian influenza viruses in poultry in China. Infection, Genetics and Evolution, 2015, 31, 64-67.	2.3	48
5	Genetic and molecular characterization of H9N2 and H5 avian influenza viruses from live poultry markets in Zhejiang Province, eastern China. Scientific Reports, 2015, 5, 17508.	3.3	40
6	Multiple amino acid substitutions involved in the adaptation of avian-origin influenza A (H10N7) virus in mice. Archives of Virology, 2016, 161, 977-980.	2.1	35
7	Analysis of the immunogenicity and bioactivities of a split influenza A/H7N9 vaccine mixed with MF59 adjuvant in BALB/c mice. Vaccine, 2016, 34, 2362-2370.	3.8	34
8	MicroRNA-181 expression regulates specific post-transcriptional level of SAMHD1 expression in vitro. Biochemical and Biophysical Research Communications, 2014, 452, 760-767.	2.1	33
9	Open-label phase I clinical trial of Ad5-EBOV in Africans in China. Human Vaccines and Immunotherapeutics, 2017, 13, 2078-2085.	3.3	28
10	Prevalence and Influencing Factors of Thyroid Dysfunction in HIV-Infected Patients. BioMed Research International, 2016, 2016, 1-11.	1.9	26
11	Correlation between reported human infection with avian influenza A H7N9 virus and cyber user awareness: what can we learn from digital epidemiology?. International Journal of Infectious Diseases, 2014, 22, 1-3.	3.3	24
12	A Decline in HIV and Syphilis Epidemics in Chinese Female Sex Workers (2000–2011): A Systematic Review and Meta-Analysis. PLoS ONE, 2013, 8, e82451.	2.5	23
13	Generation of neutralizing and non-neutralizing monoclonal antibodies against H7N9 influenza virus. Emerging Microbes and Infections, 2020, 9, 664-675.	6.5	21
14	Amino acid substitutions occurring during adaptation of an emergent H5N6 avian influenza virus to mammals. Archives of Virology, 2016, 161, 1665-1670.	2.1	20
15	Dynamic Characteristic Analysis of Antibodies in Patients With COVID-19: A 13-Month Study. Frontiers in Immunology, 2021, 12, 708184.	4.8	19
16	Comparison of a New Gold Immunochromatographic Assay for the Rapid Diagnosis of the Novel Influenza A (H7N9) Virus with Cell Culture and a Real-Time Reverse-Transcription PCR Assay. BioMed Research International, 2014, 2014, 1-6.	1.9	15
17	Amino Acid Substitutions HA A150V, PA A343T, and PB2 E627K Increase the Virulence of H5N6 Influenza Virus in Mice. Frontiers in Microbiology, 2018, 9, 453.	3.5	15
18	Development of a colloidal gold-based immunochromatographic strip test using two monoclonal antibodies to detect H7N9 avian influenza virus. Virus Genes, 2020, 56, 396-400.	1.6	14

Nanping Wu

#	Article	IF	CITATIONS
19	Molecular characterization of novel reassortant H6N2 subtype avian influenza viruses isolated from poultry in Eastern China, in 2014. Infection, Genetics and Evolution, 2015, 36, 41-45.	2.3	13
20	RbAp48, a novel inhibitory factor that regulates the transcription of human immunodeficiency virus type 1. International Journal of Molecular Medicine, 2016, 38, 267-274.	4.0	13
21	Severe dyslipidemia and immune activation in HIV patients with dysglycemia. HIV Clinical Trials, 2016, 17, 189-196.	2.0	13
22	Characterization of Novel Reassortant Influenza A (H5N2) Viruses Isolated from Poultry in Eastern China, 2015. Frontiers in Microbiology, 2017, 8, 741.	3.5	13
23	Isolation and characterization of novel reassortant H6N1 avian influenza viruses from chickens in Eastern China. Virology Journal, 2018, 15, 164.	3.4	13
24	Establishment and development of national community-based collaborative innovation demonstration areas to achieve the control target of hepatitis B in China. BMC Infectious Diseases, 2019, 19, 617.	2.9	13
25	Molecular characterization of H10 subtype avian influenza viruses isolated from poultry in Eastern China. Archives of Virology, 2019, 164, 159-179.	2.1	13
26	Novel reassortant H10N7 avian influenza viruses isolated from chickens in Eastern China. Journal of Clinical Virology, 2015, 65, 58-61.	3.1	12
27	Isolation and genetic characterization of novel reassortant H6N6 subtype avian influenza viruses isolated from chickens in eastern China. Archives of Virology, 2016, 161, 1859-1872.	2.1	12
28	Establishment of a multiplex real-time RT-PCR assay for rapid identification of H6 subtype avian influenza viruses. Archives of Virology, 2018, 163, 1671-1675.	2.1	12
29	Characterization of HIV-1 subtypes and transmitted drug resistance among treatment-naive HIV-infected individuals in Zhejiang, China, 2014-2017. Archives of Virology, 2018, 163, 2233-2237.	2.1	12
30	Molecular characterization of a reassortant H11N9 subtype avian influenza virus isolated from a domestic duck in Eastern China. Archives of Virology, 2015, 160, 2595-2601.	2.1	11
31	Elevated expression of miR-155 is associated with the differentiation of CD8+ T cells in patients with HIV-1. Molecular Medicine Reports, 2017, 16, 1584-1589.	2.4	11
32	Development of a TaqMan MGB RT-PCR assay for the detection of type A and subtype H10 avian influenza viruses. Archives of Virology, 2018, 163, 2497-2501.	2.1	11
33	Changes in Lipid Indices in HIV+ Cases on HAART. BioMed Research International, 2019, 2019, 1-8.	1.9	11
34	Genetic characterization of natural reassortant H4 subtype avian influenza viruses isolated from domestic ducks in Zhejiang province in China from 2013 to 2014. Virus Genes, 2015, 51, 347-355.	1.6	10
35	Isolation and characterization of a novel H10N2 avian influenza virus from a domestic duck in Eastern China. Infection, Genetics and Evolution, 2015, 29, 1-5.	2.3	10
36	Genetic and molecular characterization of a novel reassortant H2N8 subtype avian influenza virus isolated from a domestic duck in Zhejiang Province in China. Virus Genes, 2016, 52, 863-866.	1.6	10

NANPING WU

#	Article	IF	CITATIONS
37	Risk Factors for HIV Diagnosis Among Men Who Have Sex with Men: Results of a Case–Control Study in One Sample of Eastern China. AIDS Research and Human Retroviruses, 2016, 32, 1163-1168.	1.1	10
38	Amino acid substitutions involved in the adaptation of a novel highly pathogenic H5N2 avian influenza virus in mice. Virology Journal, 2016, 13, 159.	3.4	10
39	Molecular characterization of a novel reassortant H7N6 subtype avian influenza virus from poultry in Eastern China, in 2016. Archives of Virology, 2017, 162, 1341-1347.	2.1	10
40	High-throughput sequencing identifies HIV-1-replication- and latency-related miRNAs in CD4+ T cell lines. Archives of Virology, 2017, 162, 1933-1942.	2.1	10
41	Characterization of reassortant H1-subtype avian influenza viruses isolated from poultry in Zhejiang Province in China from 2013 to 2015. Archives of Virology, 2017, 162, 3493-3500.	2.1	10
42	Heterogeneous Evolution of HIV-1 CRF01_AE in Men Who Have Sex with Men (MSM) and Other Populations in China. PLoS ONE, 2015, 10, e0143699.	2.5	10
43	Virulence of an H5N8 highly pathogenic avian influenza is enhanced by the amino acid substitutions PB2 E627K and HA A149V. Infection, Genetics and Evolution, 2017, 54, 347-354.	2.3	9
44	hsa-miR-191-5p inhibits replication of human immunodeficiency virus type 1 by downregulating the expression of NUP50. Archives of Virology, 2021, 166, 755-766.	2.1	9
45	Preclinical evaluation of the safety and pathogenicity of a live attenuated recombinant influenza A/H7N9 seed strain and corresponding MF59-adjuvanted split vaccine. Oncotarget, 2016, 7, 81012-81025.	1.8	9
46	Near Full-Length Genome Identification of a Novel HIV-1 Recombinant Form (CRF01_AE/CRF07_BC) in Zhejiang, China. AIDS Research and Human Retroviruses, 2016, 32, 900-903.	1.1	8
47	Retinoblastoma binding protein 4 represses HIV-1 long terminal repeat–mediated transcription by recruiting NR2F1 and histone deacetylase. Acta Biochimica Et Biophysica Sinica, 2019, 51, 934-944.	2.0	7
48	Differential expression of innate immunity regulation genes in chronic HIV-1 infected adults. Cytokine, 2020, 126, 154871.	3.2	7
49	Development and evaluation of a TaqMan MGB RT-PCR assay for detection of H5 and N8 subtype influenza virus. BMC Infectious Diseases, 2020, 20, 550.	2.9	7
50	Longevity of protective immune responses induced by a split influenza A (H7N9) vaccine mixed with MF59 adjuvant in BALB/c mice. Oncotarget, 2017, 8, 91828-91840.	1.8	7
51	Gp120 binding with DC-SIGN induces reactivation of HIV-1 provirus via the NF-κB signaling pathway. Acta Biochimica Et Biophysica Sinica, 2016, 48, 275-281.	2.0	6
52	Development of a monoclonal antibodyâ€based antigen capture enzymeâ€linked immunosorbent assay for detection of H7N9 subtype avian influenza virus. Journal of Medical Virology, 2021, 93, 3939-3943.	5.0	6
53	A multiplex real-time RT-PCR method for detecting H5, H7 and H9 subtype avian influenza viruses in field and clinical samples. Virus Research, 2022, 309, 198669.	2.2	6
54	Molecular characterization of a novel reassortant H1N2 influenza virus containing genes from the 2009 pandemic human H1N1 virus in swine from eastern China. Virus Genes, 2016, 52, 405-410.	1.6	5

NANPING WU

#	Article	IF	CITATIONS
55	Identification of a Novel HIV Type 1 Recombinant Form (CRF01_AE/CRF07_BC) in Men Who Have Sex with Men in Zhejiang, China. AIDS Research and Human Retroviruses, 2017, 33, 728-734.	1.1	5
56	Molecular characterization and antigenic analysis of reassortant H9N2 subtype avian influenza viruses in Eastern China in 2016. Virus Research, 2021, 306, 198577.	2.2	5
57	The tradition algorithm approach underestimates the prevalence of serodiagnosis of syphilis in HIV-infected individuals. PLoS Neglected Tropical Diseases, 2017, 11, e0005758.	3.0	5
58	The effect of highly active antiretroviral therapy on liver function in human immunodeficiency virus-infected pediatric patients with or without hepatitis virus co-infection. Journal of Research in Medical Sciences, 2015, 20, 127-32.	0.9	5
59	Butyrylcholinesterase Levels on Admission Predict Severity and 12-Month Mortality in Hospitalized AIDS Patients. Mediators of Inflammation, 2018, 2018, 1-10.	3.0	4
60	Development of an antigen-capture enzyme-linked immunosorbent assay and immunochromatographic strip based on monoclonal antibodies for detection of H6 avian influenza viruses. Archives of Virology, 2020, 165, 1129-1139.	2.1	4
61	Development and application of a real-time RT-PCR assay to rapidly detect H2 subtype avian influenza A viruses. Journal of Veterinary Diagnostic Investigation, 2021, 33, 577-581.	1.1	4
62	RNA sequencing of CD4 T-cells reveals the relationships between lncRNA-mRNA co-expression in elite controller vs. HIV-positive infected patients. PeerJ, 2020, 8, e8911.	2.0	4
63	Preliminary research on the co-infection of human immunodeficiency virus and hepatitis virus in intravenous drug users. Chinese Medical Journal, 2003, 116, 1318-20.	2.3	4
64	Large-scale HIV testing in the older population in China: findings from a cross-sectional study. International Journal of STD and AIDS, 2014, 25, 650-655.	1.1	3
65	Isolation and molecular characterization of reassortant H11N3 subtype avian influenza viruses isolated from domestic ducks in Zhejiang Province in China. Virus Genes, 2016, 52, 732-737.	1.6	3
66	Amino acid substitutions involved in the adaptation of a novel H7N7 avian influenza virus in mice. Research in Veterinary Science, 2020, 130, 203-206.	1.9	2
67	Inactivation of Latent HIV-1 Proviral DNA Using Clustered Regularly Interspaced Short Palindromic Repeats/Cas9 Treatment and the Assessment of Off-Target Effects. Frontiers in Microbiology, 2021, 12, 629153.	3.5	2
68	Genetic analysis and biological characteristics of novel clade 2.3.4.4 reassortment H5N6 avian influenza viruses from poultry in eastern China in 2016. International Journal of Infectious Diseases, 2021, 110, 436-448.	3.3	2
69	Antigen-capture ELISA and immunochromatographic test strip to detect the H9N2 subtype avian influenza virus rapidly based on monoclonal antibodies. Virology Journal, 2021, 18, 198.	3.4	2
70	Generation, characterization, and protective ability of mouse monoclonal antibodies against the HA of A (H1N1) influenza virus. Journal of Medical Virology, 2022, 94, 2558-2567.	5.0	2
71	Isolation and molecular characterization of an H5N1 swine influenza virus in China in 2015. Archives of Virology, 2018, 163, 701-705.	2.1	1
72	Correlates of recent HIV infection among men who have sex with men recruited through the internet in Huzhou City, Eastern China. Journal of International Medical Research, 2018, 46, 5052-5061.	1.0	1

NANPING WU

#	Article	IF	CITATIONS
73	Adaptive amino acid substitutions enhance the virulence of an avian-origin H6N1 influenza virus in mice. Infection, Genetics and Evolution, 2019, 74, 103918.	2.3	1
74	Differences of cytotoxic T-lymphocyte pressure and divergent evolution of several CRF07_BC clusters circulating in men who have sex with men in China. Infection, Genetics and Evolution, 2020, 85, 104486.	2.3	1
75	Isolation and characterization of two novel reassortant H5N6 avian influenza viruses from waterfowl in eastern China. Archives of Virology, 2021, 166, 1197-1201.	2.1	1
76	Increased virulence of a novel reassortant H1N3 avian influenza virus in mice as a result of adaptive amino acid substitutions. Virus Genes, 0, , .	1.6	1
77	Development of an antigen-ELISA and a colloidal gold–based immunochromatographic strip based on monoclonal antibodies for detection of avian influenza A(H5) viruses. Journal of Veterinary Diagnostic Investigation, 2021, 33, 969-974.	1.1	0