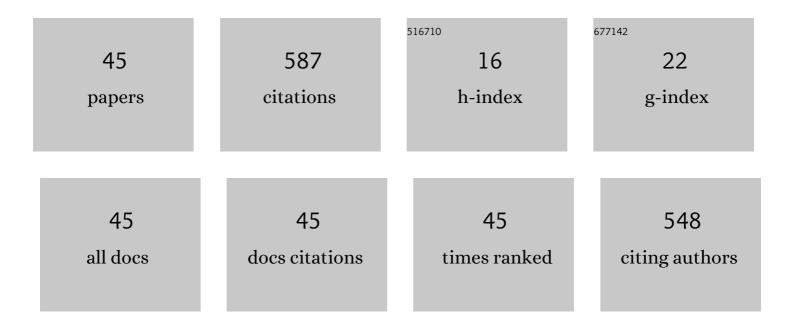
Dongmei Yan

List of Publications by Year in descending order

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DONCMELYAN

#	Article	IF	CITATIONS
1	Molecular analysis of Coxsackievirus A24 variant isolates from three outbreaks of acute hemorrhagic conjunctivitis in 1988, 1994 and 2007 in Beijing, China. Virologica Sinica, 2022, , .	3.0	1
2	Molecular Epidemiology and Evolution of Coxsackievirus A9. Viruses, 2022, 14, 822.	3.3	6
3	Whole-genome analysis of coxsackievirus B3 reflects its genetic diversity in China and worldwide. Virology Journal, 2022, 19, 69.	3.4	7
4	Identification of the first C1 subgenotype of enterovirus 71 in the Chinese mainland in a retrospective study. Virology Journal, 2022, 19, 83.	3.4	1
5	Implication of a High Risk for Type 2 Vaccine-Derived Poliovirus Emergence and Transmission After the Switch From Trivalent to Bivalent Oral Poliovirus Vaccine. Journal of Infectious Diseases, 2021, 223, 113-118.	4.0	10
6	Detection of multiple viruses potentially infecting humans in sewage water from Xinjiang Uygur Autonomous Region, China. Science of the Total Environment, 2021, 754, 142322.	8.0	11
7	Molecular epidemiological characteristics of echovirus 6 in mainland China: extensive circulation of genotype F from 2007 to 2018. Archives of Virology, 2021, 166, 1305-1312.	2.1	1
8	Coxsackievirus B4: an underestimated pathogen associated with a hand, foot, and mouth disease outbreak. Archives of Virology, 2021, 166, 2225-2234.	2.1	7
9	New Simian Enterovirus 19 (EV-A122) Strains in China Reveal Large-Scale Inter-Serotype Recombination between Simian EV-As. Virologica Sinica, 2021, 36, 1652-1655.	3.0	3
10	Monsavirus in monkey rectal swab and throat swab specimens in China: Proposal for Posaliviridae as a new family in Picornavirales. Virus Research, 2021, 303, 198501.	2.2	2
11	Circulation of Type 2 Vaccine-Derived Poliovirus in China in 2018–2019. Open Forum Infectious Diseases, 2021, 8, ofab535.	0.9	5
12	Development of a real-time RT-PCR assay for the detection of pan-human parechoviruses. Virology Journal, 2021, 18, 227.	3.4	2
13	Immunogenicity of Oral Polio Vaccine and Salk Inactive Polio Vaccine Against Xinjiang Imported Type 1 Wild Poliovirus. Clinical Infectious Diseases, 2020, 70, 1980-1984.	5.8	3
14	Genetic Diversity Analysis of Coxsackievirus A8 Circulating in China and Worldwide Reveals a Highly Divergent Genotype. Viruses, 2020, 12, 1061.	3.3	4
15	Global Spread of the B5 Subgenotype EV-A71 and the Phylogeographical Analysis of Chinese Migration Events. Frontiers in Cellular and Infection Microbiology, 2020, 10, 475.	3.9	6
16	A novel interspecies recombinant enterovirus (Enterovirus A120) isolated from a case of acute flaccid paralysis in China. Emerging Microbes and Infections, 2020, 9, 1733-1743.	6.5	6
17	Excretion of SARS-CoV-2 through faecal specimens. Emerging Microbes and Infections, 2020, 9, 2501-2508.	6.5	45
18	Molecular typing and characterization of a novel genotype of EV-B93 isolated from Tibet, China. PLoS ONE, 2020, 15, e0237652.	2.5	2

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19	The Husavirus Posa-Like Viruses in China, and a New Group of Picornavirales. Viruses, 2020, 12, 995.	3.3	8
20	Genetic recombination in fast-spreading coxsackievirus A6 variants: a potential role in evolution and pathogenicity. Virus Evolution, 2020, 6, veaa048.	4.9	13
21	Phylogenetic characteristics and molecular epidemiological analysis of novel enterovirus EV-B83 isolated from Tibet, China. Scientific Reports, 2020, 10, 6630.	3.3	9
22	Molecular Epidemiological, Serological, and Pathogenic Analysis of EV-B75 Associated With Acute Flaccid Paralysis Cases in Tibet, China. Frontiers in Microbiology, 2020, 11, 632552.	3.5	1
23	Genomic epidemiology of coxsackievirus A16 in mainland of China, 2000–18. Virus Evolution, 2020, 6, veaa084.	4.9	21
24	Antibody Response to COVID-19 Virus — Heilongjiang Province and Gansu Province, China, 2020. China CDC Weekly, 2020, 2, 645-650.	2.3	0
25	Multiple genotypes of Echovirus 11 circulated in mainland China between 1994 and 2017. Scientific Reports, 2019, 9, 10583.	3.3	14
26	Emerging recombination of the C2 sub-genotype of HFMD-associated CV-A4 is persistently and extensively circulating in China. Scientific Reports, 2019, 9, 13668.	3.3	7
27	Two Coxsackievirus B3 outbreaks associated with hand, foot, and mouth disease in China and the evolutionary history worldwide. BMC Infectious Diseases, 2019, 19, 466.	2.9	23
28	Phylogenetic analysis and phenotypic characterisatics of two Tibet EV-C96 strains. Virology Journal, 2019, 16, 40.	3.4	4
29	Genetic characterization and molecular epidemiological analysis of novel enterovirus EV-B80 in China. Emerging Microbes and Infections, 2018, 7, 1-12.	6.5	23
30	Persistent circulation of genotype D coxsackievirus A2 in mainland of China since 2008. PLoS ONE, 2018, 13, e0204359.	2.5	11
31	The emerging sub-genotype C2 of CoxsackievirusA10 Associated with Hand, Foot and Mouth Disease extensively circulating in mainland of China. Scientific Reports, 2018, 8, 13357.	3.3	24
32	Antigenic characteristics and genomic analysis of novel EV-A90 enteroviruses isolated in Xinjiang, China. Scientific Reports, 2018, 8, 10247.	3.3	9
33	Phylogenetic Characterizations of Highly Mutated EV-B106 Recombinants Showing Extensive Genetic Exchanges with Other EV-B in Xinjiang, China. Scientific Reports, 2017, 7, 43080.	3.3	16
34	Persistent circulation of Coxsackievirus A6 of genotype D3 in mainland of China between 2008 and 2015. Scientific Reports, 2017, 7, 5491.	3.3	66
35	lsolation of an imported subgenotype B5 strain of human enterovirus A71 in Chongqing City, China, 2014. Virology Journal, 2016, 13, 115.	3.4	7
36	Two Genotypes of Coxsackievirus A2 Associated with Hand, Foot, and Mouth Disease Circulating in China since 2008. PLoS ONE, 2016, 11, e0169021.	2.5	17

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37	Circulation of multiple serotypes of highly divergent enterovirus C in the Xinjiang Uighur Autonomous Region of China. Scientific Reports, 2016, 6, 33595.	3.3	18
38	A Novel Recombinant Enterovirus Type EV-A89 with Low Epidemic Strength in Xinjiang, China. Scientific Reports, 2015, 5, 18558.	3.3	19
39	An Insight into Recombination with Enterovirus Species C and Nucleotide G-480 Reversion from the Viewpoint of Neurovirulence of Vaccine-Derived Polioviruses. Scientific Reports, 2015, 5, 17291.	3.3	25
40	Molecular typing and characterization of a new serotype of human enterovirus (EV-B111) identified in China. Virus Research, 2014, 183, 75-80.	2.2	21
41	Limited and Localized Outbreak of Newly Emergent Type 2 Vaccine-Derived Poliovirus in Sichuan, China. Vaccine Journal, 2014, 21, 1012-1018.	3.1	17
42	Phylogenetic evidence for multiple intertypic recombinations in enterovirus B81 strains isolated in Tibet, China. Scientific Reports, 2014, 4, 6035.	3.3	23
43	Isolation and Characterization of a Type 2 Vaccine-Derived Poliovirus from Environmental Surveillance in China, 2012. PLoS ONE, 2013, 8, e83975.	2.5	19
44	Emergence and Localized Circulation of a Vaccine-Derived Poliovirus in an Isolated Mountain Community in Guangxi, China. Journal of Clinical Microbiology, 2010, 48, 3274-3280.	3.9	25
45	Outbreak of acute hemorrhagic conjunctivitis in Yunnan, People's Republic of China, 2007. Virology Journal, 2010, 7, 138.	3.4	25