Emergence of ON1 genotype of human respiratory sync between 2011 and 2015

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Citation Report

#	Article	IF	Citations
1	Emergence of BA9 genotype of human respiratory syncytial virus subgroup B in China from 2006 to 2014. Scientific Reports, 2017, 7, 16765.	1.6	24
2	Sequence Analysis of the Fusion Protein Gene of Human Respiratory Syncytial Virus Circulating in China from 2003 to 2014. Scientific Reports, 2018, 8, 17618.	1.6	5
3	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.	1.6	24
4	Virus isolation and genotype identification of human respiratory syncytial virus in Guizhou Province, China. Brazilian Journal of Infectious Diseases, 2019, 23, 427-434.	0.3	2
5	Epidemiological characteristics and phylogenic analysis of human respiratory syncytial virus in patients with respiratory infections during 2011â€“2016 in southern China. International Journal of Infectious Diseases, 2020, 90, 5-17.	1.5	22
6	The emergence of subgenotype ONâ€1 of Human orthopneumovirus type A in Riyadh, Saudi Arabia: A new episode of the virus epidemiological dynamic. Journal of Medical Virology, 2020, 92, 1133-1140.	2.5	4
7	Epidemiological characteristics of respiratory viruses in patients with acute respiratory infections during 2009â€“2018 in southern China. International Journal of Infectious Diseases, 2020, 98, 21-32.	1.5	12
8	Emerging Human Metapneumovirus Gene Duplication Variants in Patients with Severe Acute Respiratory Infection, China, 2017â€“2019. Emerging Infectious Diseases, 2021, 27, 275-277.	2.0	11
10	Evolutionary dynamics of group A and B respiratory syncytial virus in China, 2009-2018. Archives of Virology, 2021, 166, 2407-2418.	0.9	5
11	A multi-center study on Molecular Epidemiology of Human Respiratory Syncytial Virus from Children with Acute Lower Respiratory Tract Infections in the Mainland of China between 2015 and 2019. Virologica Sinica, 2021, 36, 1475-1483.	1.2	12
12	Human Respiratory Syncytial Virus Detected in Mountain Gorilla Respiratory Outbreaks. EcoHealth, 2020, 17, 449-460.	0.9	19
13	Sequence analysis of the G gene of hRSVA ON1 genotype from Egyptian children with acute respiratory tract infections. Journal of Medical Microbiology, 2018, 67, 387-391.	0.7	4
14	Molecular characterization of respiratory syncytial viruses circulating in a paediatric cohort in Amman, Jordan. Microbial Genomics, 2021, 7, .	1.0	8
15	Molecular epidemiology of human respiratory syncytial virus among children in Japan during three seasons and hospitalization risk of genotype ON1. PLoS ONE, 2018, 13, e0192085.	1.1	29
16	Genetic characterization of G protein in respiratory syncytial virus ON-1 genotype in Tehran. Future Virology, 2020, 15, 725-734.	0.9	1
17	Molecular Evolution of Attachment Glycoprotein (G) and Fusion Protein (F) Genes of Respiratory Syncytial Virus ON1 and BA9 Strains in Xiamen, China. Microbiology Spectrum, 2022, 10, e0208321.	1.2	6
18	Genetic diversity and epidemiological features of respiratory syncytial virus, Beijing, 2015â€“2019: A multicenter and all-age groups study. Journal of Infection, 2022, 85, 75-85.	1.7	7
19	The A2c _{111ntâ€dup} Variants of Human Metapneumovirus Predominantly Circulating in Qingdao, China, during 2018 and 2019. Journal of Medical Virology, 0, , .	2.5	0

#	Article	IF	Citations
20	Resurgence of Respiratory Syncytial Virus Infection During COVID-19 Pandemic Among Children in Shanghai, China. Frontiers in Microbiology, 0, 13, .	1.5	30
21	Genome analysis of human respiratory syncytial virus in Fujian Province, Southeast China. Infection, Genetics and Evolution, 2022, 103, 105329.	1.0	3
22	Clinical characteristics and molecular epidemiology of human metapneumovirus in children with acute lower respiratory tract infections in China, 2017 to 2019: A multicentre prospective observational study. Virologica Sinica, 2022, 37, 874-882.	1.2	5
23	Changes in endemic patterns of respiratory syncytial virus infection in pediatric patients under the pressure of nonpharmaceutical interventions for COVIDâ€19 in Beijing, China. Journal of Medical Virology, 2023, 95, .	2.5	9
25	Circulation pattern and genetic variation of human respiratory syncytial virus in China during 2008â€“2021. Journal of Medical Virology, 2023, 95, .	2.5	3