Lin-Qing Zhao

List of Publications by Year in descending order

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17	202	1040056	940533
17	292	9	16
papers	citations	h-index	g-index
21	21	21	476
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Genetic Variation in Attachment Glycoprotein Genes of Human Respiratory Syncytial Virus Subgroups A and B in Children in Recent Five Consecutive Years. PLoS ONE, 2013, 8, e75020.	2.5	85
2	Detection of a Bocavirus Circular Genome in Fecal Specimens from Children with Acute Diarrhea in Beijing, China. PLoS ONE, 2012, 7, e48980.	2.5	41
3	Risk of acute gastroenteritis associated with human bocavirus infection in children: A systematic review and meta-analysis. PLoS ONE, 2017, 12, e0184833.	2.5	34
4	Four DNA extraction methods used in loop-mediated isothermal amplification for rapid adenovirus detection. Journal of Virological Methods, 2014, 204, 49-52.	2.1	19
5	Early Serum HBsAg Kinetics as Predictor of HBsAg Loss in Patients with HBeAg-Negative Chronic Hepatitis B after Treatment with Pegylated Interferonα-2a. Virologica Sinica, 2021, 36, 311-320.	3.0	17
6	Prevalence analysis of different human bocavirus genotypes in pediatric patients revealed intra-genotype recombination. Infection, Genetics and Evolution, 2014, 27, 382-388.	2.3	16
7	Surges of hospital-based rhinovirus infection during the 2020 coronavirus disease-19 (COVID-19) pandemic in Beijing, China. World Journal of Pediatrics, 2021, 17, 590-596.	1.8	14
8	Tracing the emerging genotypes of human respiratory syncytial virus in Beijing by evolution analysis of the attachment glycoprotein (G) gene. Infection, Genetics and Evolution, 2018, 65, 18-27.	2.3	12
9	Association of Different Human Rhinovirus Species with Asthma in Children. Chinese Medical Journal, 2016, 129, 1513-1518.	2.3	11
10	Consolidation treatment needed for sustained HBsAg-negative response induced by interferon-alpha in HBeAg positive chronic hepatitis B patients. Virologica Sinica, 2022, 37, 390-397.	3.0	10
11	Reverse transcription genome exponential amplification reaction assay for rapid and universal detection of human rhinoviruses. Archives of Virology, 2016, 161, 1891-1898.	2.1	6
12	Application of Human Adenovirus Genotyping by Phylogenetic Analysis in an Outbreak to Identify Nosocomial Infection. Virologica Sinica, 2021, 36, 393-401.	3.0	5
13	Virus profile in children with acute respiratory infections with various severities in Beijing, China. Chinese Medical Journal, 2014, 127, 3706-11.	2.3	5
14	Prevalence and Phylogenetic Analysis of Human Bocaviruses 1-4 in Pediatric Patients with Various Infectious Diseases. PLoS ONE, 2016, 11 , e0160603.	2.5	4
15	Multiple RNA virus matrix proteins interact with SLD5 to manipulate host cell cycle. Journal of General Virology, 2021, 102, .	2.9	1
16	Human rhinovirus C infection is associated with asthma in children determined by xTAG respiratory viral panel FAST. Virologica Sinica, 2017, 32, 171-174.	3.0	0
17	Keeping all secondary structures of the non-coding region in the circular genome of human bocavirus 2 is important for DNA replication and virus assembly, as revealed by three hetero-recombinant genomic clones. Emerging Microbes and Infections, 2019, 8, 1563-1573.	6.5	0