Lu Lu

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6860117/publications.pdf

Version: 2024-02-01

18	1,066	11	19
papers	citations	h-index	g-index
23	23	23	2090
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Role for migratory wild birds in the global spread of avian influenza H5N8. Science, 2016, 354, 213-217.	12.6	362
2	Monitoring SARS-CoV-2 Circulation and Diversity through Community Wastewater Sequencing, the Netherlands and Belgium. Emerging Infectious Diseases, 2021, 27, 1405-1415.	4.3	168
3	Methane oxidation coupled to oxygenic photosynthesis in anoxic waters. ISME Journal, 2015, 9, 1991-2002.	9.8	135
4	Adaptation, spread and transmission of SARS-CoV-2 in farmed minks and associated humans in the Netherlands. Nature Communications, 2021, 12, 6802.	12.8	81
5	Reassortment patterns of avian influenza virus internal segments among different subtypes. BMC Evolutionary Biology, 2014, 14, 16.	3.2	77
6	Novel genetic reassortants in H9N2 influenza A viruses and their diverse pathogenicity to mice. Virology Journal, 2011, 8, 505.	3.4	44
7	Quantifying predictors for the spatial diffusion of avian influenza virus in China. BMC Evolutionary Biology, 2017, 17, 16.	3.2	31
8	Hepatitis C virus infection induces the expression of amphiregulin, a factor related to the activation of cellular survival pathways and required for efficient viral assembly. Journal of General Virology, 2011, 92, 2237-2248.	2.9	27
9	Determining the Phylogenetic and Phylogeographic Origin of Highly Pathogenic Avian Influenza (H7N3) in Mexico. PLoS ONE, 2014, 9, e107330.	2.5	25
10	Genetic diversity and cross-species transmission of kobuviruses in Vietnam. Virus Evolution, 2018, 4, vey002.	4.9	18
11	Re-emergence of H5N8 highly pathogenic avian influenza virus in wild birds, China. Emerging Microbes and Infections, 2021, 10, 1819-1823.	6.5	17
12	Segmentation and shielding of the most vulnerable members of the population as elements of an exit strategy from COVID-19 lockdown. Philosophical Transactions of the Royal Society B: Biological Sciences, 2021, 376, 20200275.	4.0	15
13	What are SARS-CoV-2 genomes from the WHO Africa region member states telling us?. BMJ Global Health, 2021, 6, e004408.	4.7	9
14	Genetic correlation between current circulating H1N1 swine and human influenza viruses. Journal of Clinical Virology, 2010, 49, 186-191.	3.1	8
15	Evolution and phylogeographic dissemination of endemic porcine picornaviruses in Vietnam. Virus Evolution, 2016, 2, vew001.	4.9	7
16	Epidemiology and Phylogenetic Analysis of Viral Respiratory Infections in Vietnam. Frontiers in Microbiology, 2020, 11, 833.	3.5	6
17	No Exchange of Picornaviruses in Vietnam between Humans and Animals in a High-Risk Cohort with Close Contact despite High Prevalence and Diversity. Viruses, 2021, 13, 1709.	3.3	2
18	Dealing with Highly Pathogenic Avian Influenza: An Impending Crisis. Innovation(China), 2021, 2, 100084.	9.1	1