Huaiyu Tian

List of Publications by Year in descending order

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331259 288905 5,821 41 21 40 h-index citations g-index papers 49 49 49 9287 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	The effect of human mobility and control measures on the COVID-19 epidemic in China. Science, 2020, 368, 493-497.	6.0	2,168
2	An investigation of transmission control measures during the first 50 days of the COVID-19 epidemic in China. Science, 2020, 368, 638-642.	6.0	1,554
3	Reduction of secondary transmission of SARS-CoV-2 in households by face mask use, disinfection and social distancing: a cohort study in Beijing, China. BMJ Global Health, 2020, 5, e002794.	2.0	382
4	Crowding and the shape of COVID-19 epidemics. Nature Medicine, 2020, 26, 1829-1834.	15.2	204
5	Modelling COVID-19. Nature Reviews Physics, 2020, 2, 279-281.	11.9	174
6	Avian influenza H5N1 viral and bird migration networks in Asia. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 172-177.	3.3	169
7	Human Infection with Influenza Virus A(H10N8) from Live Poultry Markets, China, 2014. Emerging Infectious Diseases, 2014, 20, 2076-9.	2.0	94
8	Increasing airline travel may facilitate co-circulation of multiple dengue virus serotypes in Asia. PLoS Neglected Tropical Diseases, 2017, 11, e0005694.	1.3	86
9	Urbanization prolongs hantavirus epidemics in cities. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 4707-4712.	3.3	72
10	Interannual cycles of Hantaan virus outbreaks at the human–animal interface in Central China are controlled by temperature and rainfall. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 8041-8046.	3.3	67
11	The ecological dynamics of hantavirus diseases: From environmental variability to disease prevention largely based on data from China. PLoS Neglected Tropical Diseases, 2019, 13, e0006901.	1.3	58
12	Normalized difference vegetation index dynamic and spatiotemporal distribution of migratory birds in the Poyang Lake wetland, China. Ecological Indicators, 2014, 47, 219-230.	2.6	57
13	Impact of global change on transmission of human infectious diseases. Science China Earth Sciences, 2014, 57, 189-203.	2.3	57
14	Assessing the role of live poultry trade in community-structured transmission of avian influenza in China. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 5949-5954.	3.3	43
15	Anthropogenically driven environmental changes shift the ecological dynamics of hemorrhagic fever with renal syndrome. PLoS Pathogens, 2017, 13, e1006198.	2.1	41
16	Impacts of Road Traffic Network and Socioeconomic Factors on the Diffusion of 2009 Pandemic Influenza A ($H1N1$) in Mainland China. International Journal of Environmental Research and Public Health, 2019, 16, 1223.	1.2	35
17	Associations between changes in population mobility in response to the COVID-19 pandemic and socioeconomic factors at the city level in China and country level worldwide: a retrospective, observational study. The Lancet Digital Health, 2021, 3, e349-e359.	5.9	35
18	Association between coronavirus disease 2019 (COVID-19) and long-term exposure to air pollution: Evidence from the first epidemic wave in China. Environmental Pollution, 2021, 276, 116682.	3.7	33

#	Article	IF	CITATIONS
19	Surface water areas significantly impacted 2014 dengue outbreaks in Guangzhou, China. Environmental Research, 2016, 150, 299-305.	3.7	29
20	Ecology and geography of hemorrhagic fever with renal syndrome in Changsha, China. BMC Infectious Diseases, 2013, 13, 305.	1.3	27
21	Landscape and rodent community composition are associated with risk of hemorrhagic fever with renal syndrome in two cities in China, 2006–2013. BMC Infectious Diseases, 2018, 18, 37.	1.3	24
22	Climate change suggests a shift of H5N1 risk in migratory birds. Ecological Modelling, 2015, 306, 6-15.	1.2	23
23	Migratory Whooper Swans Cygnus cygnus Transmit H5N1 Virus between China and Mongolia: Combination Evidence from Satellite Tracking and Phylogenetics Analysis. Scientific Reports, 2018, 8, 7049.	1.6	23
24	Analysis of an Outbreak of Hemorrhagic Fever with Renal Syndrome in College Students in Xi'an, China. Viruses, 2014, 6, 507-515.	1.5	22
25	Spatial heterogeneity of hemorrhagic fever with renal syndrome is driven by environmental factors and rodent community composition. PLoS Neglected Tropical Diseases, 2018, 12, e0006881.	1.3	20
26	Contrasting effects of host species and phylogenetic diversity on the occurrence of HPAI H5N1 in European wild birds. Journal of Animal Ecology, 2019, 88, 1044-1053.	1.3	20
27	Spatial, temporal and genetic dynamics of highly pathogenic avian influenza A (H5N1) virus in China. BMC Infectious Diseases, 2015, 15, 54.	1.3	19
28	A follow-up study shows that recovered patients with re-positive PCR test in Wuhan may not be infectious. BMC Medicine, 2021, $19,77$.	2.3	19
29	Time-Specific Ecologic Niche Models Forecast the Risk of Hemorrhagic Fever with Renal Syndrome in Dongting Lake District, China, 2005–2010. PLoS ONE, 2014, 9, e106839.	1.1	15
30	Intrinsic and extrinsic drivers of transmission dynamics of hemorrhagic fever with renal syndrome caused by Seoul hantavirus. PLoS Neglected Tropical Diseases, 2019, 13, e0007757.	1.3	15
31	Reâ€emerging of rabies in Shaanxi province, China, from 2009 to 2015. Journal of Medical Virology, 2017, 89, 1511-1519.	2.5	13
32	New evidence for the east–west spread of the highly pathogenic avian influenza H5N1 virus between Central Asian and east Asian-Australasian flyways in China. Emerging Microbes and Infections, 2019, 8, 823-826.	3.0	11
33	Evaluating the effectiveness of control measures in multiple regions during the early phase of the COVID-19 pandemic in 2020. Biosafety and Health, 2021, 3, 264-275.	1.2	11
34	Assessing the impact of COVID-19 border restrictions on dengue transmission in Yunnan Province, China: an observational epidemiological and phylogenetic analysis. The Lancet Regional Health - Western Pacific, 2021, 14, 100259.	1.3	11
35	Persistence and transmission of avian influenza A (H5N1): virus movement, risk factors and pandemic potential. Annals of GIS, 2015, 21, 55-68.	1.4	8
36	Genetic evidence for avian influenza H5N1 viral transmission along the Black Sea–Mediterranean Flyway. Journal of General Virology, 2016, 97, 2129-2134.	1.3	8

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37	Using Satellite Data for the Characterization of Local Animal Reservoir Populations of Hantaan Virus on the Weihe Plain, China. Remote Sensing, 2017, 9, 1076.	1.8	7
38	Risk analysis of H5N1 highly pathogenic avian influenza in poultry at the Poyang Lake area, China. Environmental Earth Sciences, 2016, 75, 1.	1.3	4
39	Population serology for SARS-CoV-2 is essential to regional and global preparedness. Lancet Microbe, The, 2020, 1, e94.	3.4	4
40	The ecological dynamics of the coronavirus epidemics during transmission from outside sources when R O is successfully managed below one. Royal Society Open Science, 2021, 8, 202234.	1.1	2
41	Anthropogenic factors and societal response to challenges in the transmission of highly pathogenic avian influenza A (H5N1). Annals of GIS, 2015, 21, 149-156.	1.4	1