Jonathan H Epstein

List of Publications by Year
in descending order

51 papers	8,277 citations	126907 33 h-index	189892 50 g-index
56 all docs	56 docs citations	56 times ranked	9706 citing authors

#	Article	IF	Citations
1	Nipah Virus Detection at Bat Roosts after Spillover Events, Bangladesh, 2012â€“2019. Emerging Infectious Diseases, 2022, 28, 1384-1392.	4.3	3
2	Functional Analysis of the Fusion and Attachment Glycoproteins of Mojiang Henipavirus. Viruses, 2021, 13, 517.	3.3	15
3	Microbicidal actives with virucidal efficacy against SARS-CoV-2 and other beta- and alpha-coronaviruses and implications for future emerging coronaviruses and other enveloped viruses. Scientific Reports, 2021, 11, 5626.	3.3	45
4	Overview of Bat and Wildlife Coronavirus Surveillance in Africa: A Framework for Global Investigations. Viruses, 2021, 13, 936.	3.3	23
5	Seasonality of Date Palm Sap Feeding Behavior by Bats in Bangladesh. EcoHealth, 2021, 18, 359-371.	2.0	2
6	Seasonal shedding patterns of diverse henipavirus-related paramyxoviruses in Egyptian rousette bats. Scientific Reports, 2021, 11, 24262.	3.3	10
7	Population genetics of fruit bat reservoir informs the dynamics, distribution and diversity of Nipah virus. Molecular Ecology, 2020, 29, 970-985.	3.9	24
8	Molecular characterization of group A rotavirus from rhesus macaques (Macaca mulatta) at humanâ€“wildlife interfaces in Bangladesh. Transboundary and Emerging Diseases, 2020, 67, 956-966.	3.0	17
9	Possibility for reverse zoonotic transmission of SARS-CoV-2 to free-ranging wildlife: A case study of bats. PLoS Pathogens, 2020, 16, e1008758.	4.7	127
10	Epidemiology and Molecular Characterization of Rotavirus A in Fruit Bats in Bangladesh. EcoHealth, 2020, 17, 398-405.	2.0	9
11	Origin and cross-species transmission of bat coronaviruses in China. Nature Communications, 2020, 11, 4235.	12.8	264
12	No Evidence of Coronaviruses or Other Potentially Zoonotic Viruses in Sunda pangolins (Manis) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 30	2.0	53
13	Nipah virus dynamics in bats and implications for spillover to humans. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 29190-29201.	7.1	119
14	Exploring the Mental Model of Cattle Farmers in Disease Prevention and Control Practices. Veterinary Sciences, 2020, 7, 27.	1.7	8
15	Pteropus lylei primarily forages in residential areas in Kandal, Cambodia. Ecology and Evolution, 2019, 9, 4181-4191.	1.9	17
16	A viral metagenomic survey identifies known and novel mammalian viruses in bats from Saudi Arabia. PLoS ONE, 2019, 14, e0214227.	2.5	36
17	Isolation and Full-Genome Characterization of Nipah Viruses from Bats, Bangladesh. Emerging Infectious Diseases, 2019, 25, 166-170.	4.3	32
18	Emerging Diseases in Bats. , 2019, , 274-279.		2

#	Article	IF	Citations
19	Food for contagion: synthesis and future directions for studying hostâ€“parasite responses to resource shifts in anthropogenic environments. Philosophical Transactions of the Royal Society B: Biological Sciences, 2018, 373, 20170102.	4.0	54
20	Convergence of Humans, Bats, Trees, and Culture in Nipah Virus Transmission, Bangladesh. Emerging Infectious Diseases, 2017, 23, 1446-1453.	4.3	76
21	Increased Morbidity and Mortality in Domestic Animals Eating Dropped and Bitten Fruit in Bangladeshi Villages: Implications for Zoonotic Disease Transmission. EcoHealth, 2016, 13, 39-48.	2.0	10
22	Molecular evidence of Ebola Reston virus infection in Philippine bats. Virology Journal, 2015, 12, 107.	3.4	71
23	Diversity of coronavirus in bats from Eastern Thailand. Virology Journal, 2015, 12, 57.	3.4	70
24	Ecological dynamics of emerging bat virus spillover. Proceedings of the Royal Society B: Biological Sciences, 2015, 282, 20142124.	2.6	375
25	Evidence for Retrovirus and Paramyxovirus Infection of Multiple Bat Species in China. Viruses, 2014, 6, 2138-2154.	3.3	25
26	Serological Evidence of Henipavirus Exposure in Cattle, Goats and Pigs in Bangladesh. PLoS Neglected Tropical Diseases, 2014, 8, e3302.	3.0	57
27	Roosting behaviour and habitat selection of Pteropus giganteus reveal potential links to Nipah virus epidemiology. Journal of Applied Ecology, 2014, 51, 376-387.	4.0	58
28	The Role of Landscape Composition and Configuration on Pteropus giganteus Roosting Ecology and Nipah Virus Spillover Risk in Bangladesh. American Journal of Tropical Medicine and Hygiene, 2014, 90, 247-255.	1.4	62
29	Isolation and characterization of a bat SARS-like coronavirus that uses the ACE2 receptor. Nature, 2013, 503, 535-538.	27.8	1,439
30	Isolation of Salmonella Virchow from a Fruit Bat (Pteropus giganteus). EcoHealth, 2013, 10, 348-351.	2.0	16
31	Interdisciplinary approaches to understanding disease emergence: The past, present, and future drivers of Nipah virus emergence. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 3681-3688.	7.1	128
32	A Strategy To Estimate Unknown Viral Diversity in Mammals. MBio, 2013, 4, e00598-13.	4.1	320
33	Ebola Virus Antibodies in Fruit Bats, Bangladesh. Emerging Infectious Diseases, 2013, 19, 270-273.	4.3	129
34	Middle East Respiratory Syndrome Coronavirus in Bats, Saudi Arabia. Emerging Infectious Diseases, 2013, 19, 1819-23.	4.3	562
35	Bats are a major natural reservoir for hepaciviruses and pegiviruses. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 8194-8199.	7.1	251
36	Risk Factors for Nipah Virus Infection among Pteropid Bats, Peninsular Malaysia. Emerging Infectious Diseases, 2013, 19, 51-60.	4.3	44

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37	Group C Betacoronavirus in Bat Guano Fertilizer, Thailand. Emerging Infectious Diseases, 2013, 19, 1349-51.	4.3	65
38	Duration of Maternal Antibodies against Canine Distemper Virus and Hendra Virus in Pteropid Bats. PLoS ONE, 2013, 8, e67584.	2.5	37
39	Agricultural intensification, priming for persistence and the emergence of Nipah virus: a lethal bat-borne zoonosis. Journal of the Royal Society Interface, 2012, 9, 89-101.	3.4	245
40	A framework for the study of zoonotic disease emergence and its drivers: spillover of bat pathogens as a case study. Philosophical Transactions of the Royal Society B: Biological Sciences, 2012, 367, 2881-2892.	4.0	156
41	Pteropid Bats are Confirmed as the Reservoir Hosts of Henipaviruses: A Comprehensive Experimental Study of Virus Transmission. American Journal of Tropical Medicine and Hygiene, 2011, 85, 946-951.	1.4	337
42	Comparison of Intravenous Medetomidine and Medetomidine/Ketamine for Immobilization of Free-Ranging Variable Flying Foxes (Pteropus hypomelanus). PLoS ONE, 2011, 6, e25361.	2.5	11
43	Identification of GBV-D, a Novel GB-like Flavivirus from Old World Frugivorous Bats (Pteropus) Tj ETQq1 1 0.784314 rgBT /Overlock 10	4.7	69
44	Pteropus vampyrus, a hunted migratory species with a multinational homeâ€range and a need for regional management. Journal of Applied Ecology, 2009, 46, 991-1002.	4.0	145
45	The Significant but Understudied Impact of Pathogen Transmission from Humans to Animals. Mount Sinai Journal of Medicine, 2009, 76, 448-455.	1.9	43
46	HenipavirusInfection in Fruit Bats (Pteropus giganteus), India. Emerging Infectious Diseases, 2008, 14, 1309-1311.	4.3	121
47	Emerging Viruses: Coming in on a Wrinkled Wing and a Prayer. Clinical Infectious Diseases, 2007, 44, 711-717.	5.8	94
48	Emerging henipaviruses and flying foxes â€“ Conservation and management perspectives. Biological Conservation, 2006, 131, 211-220.	4.1	43
49	Nipah virus: Impact, origins, and causes of emergence. Current Infectious Disease Reports, 2006, 8, 59-65.	3.0	182
50	Bats Are Natural Reservoirs of SARS-Like Coronaviruses. Science, 2005, 310, 676-679.	12.6	2,130
51	Bringing Conservation Medicine into the Veterinary Curriculum: The Tufts Example. EcoHealth, 2004, 1, S43.	2.0	7