

Peter Daszak

List of Publications by Year in descending order

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Version: 2024-02-01

301
papers

45,329
citations

3325

91
h-index

2171

202
g-index

318
all docs

318
docs citations

318
times ranked

36748
citing authors

#	ARTICLE	IF	CITATIONS
1	Healthy planet healthy people. <i>Conservation Letters</i> , 2022, 15, .	2.8	3
2	The costs and benefits of primary prevention of zoonotic pandemics. <i>Science Advances</i> , 2022, 8, eabl4183.	4.7	99
3	Investing to Both Prevent and Prepare for COVID-XX. <i>EcoHealth</i> , 2022, , 1.	0.9	2
4	Behavioral biological surveillance of emerging infectious diseases among a dynamic cohort in Thailand. <i>BMC Infectious Diseases</i> , 2022, 22, 472.	1.3	0
5	Knowledge, attitudes, and practices associated with zoonotic disease transmission risk in North Sulawesi, Indonesia. <i>One Health Outlook</i> , 2022, 4, .	1.4	3
6	Nipah Virus Detection at Bat Roosts after Spillover Events, Bangladesh, 2012–2019. <i>Emerging Infectious Diseases</i> , 2022, 28, 1384-1392.	2.0	3
7	Classification of new morbillivirus and jeilongvirus sequences from bats sampled in Brazil and Malaysia. <i>Archives of Virology</i> , 2022, 167, 1977-1987.	0.9	11
8	Lessons from COVID-19 to Help Prevent Future Pandemics. <i>China CDC Weekly</i> , 2021, 3, 132-133.	1.0	1
9	Infectious Disease Threats: A Rebound To Resilience. <i>Health Affairs</i> , 2021, 40, 204-211.	2.5	50
10	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. <i>Scientific Reports</i> , 2021, 11, 5626.	1.6	45
11	Ranking the risk of animal-to-human spillover for newly discovered viruses. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	3.3	140
12	Socializing One Health: an innovative strategy to investigate social and behavioral risks of emerging viral threats. <i>One Health Outlook</i> , 2021, 3, 11.	1.4	18
13	Science, not speculation, is essential to determine how SARS-CoV-2 reached humans. <i>Lancet, The</i> , 2021, 398, 209-211.	6.3	18
14	A Novel Potentially Recombinant Rodent Coronavirus with a Polybasic Cleavage Site in the Spike Protein. <i>Journal of Virology</i> , 2021, 95, e0117321.	1.5	16
15	Wild animal and zoonotic disease risk management and regulation in China: Examining gaps and One Health opportunities in scope, mandates, and monitoring systems. <i>One Health</i> , 2021, 13, 100301.	1.5	18
16	Decoding the RNA viromes in rodent lungs provides new insight into the origin and evolutionary patterns of rodent-borne pathogens in Mainland Southeast Asia. <i>Microbiome</i> , 2021, 9, 18.	4.9	43
17	Seasonality of Date Palm Sap Feeding Behavior by Bats in Bangladesh. <i>EcoHealth</i> , 2021, 18, 359-371.	0.9	2
18	Environmental Change and Zoonotic Disease Risk at Human-Macaque Interfaces in Bangladesh. <i>EcoHealth</i> , 2021, 18, 487-499.	0.9	2

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19	Protection of wetlands as a strategy for reducing the spread of avian influenza from migratory waterfowl. <i>Ambio</i> , 2020, 49, 939-949.	2.8	9
20	Population genetics of fruit bat reservoir informs the dynamics, distribution and diversity of Nipah virus. <i>Molecular Ecology</i> , 2020, 29, 970-985.	2.0	24
21	Characterizing and quantifying the wildlife trade network in Sulawesi, Indonesia. <i>Global Ecology and Conservation</i> , 2020, 21, e00887.	1.0	23
22	Ecology and economics for pandemic prevention. <i>Science</i> , 2020, 369, 379-381.	6.0	411
23	Was the COVID-19 pandemic avoidable? A call for a "resolution-oriented" approach in pathogen evolutionary ecology to prevent future outbreaks. <i>Ecology Letters</i> , 2020, 23, 1557-1560.	3.0	27
24	Possibility for reverse zoonotic transmission of SARS-CoV-2 to free-ranging wildlife: A case study of bats. <i>PLoS Pathogens</i> , 2020, 16, e1008758.	2.1	127
25	Epidemiology and Molecular Characterization of Rotavirus A in Fruit Bats in Bangladesh. <i>EcoHealth</i> , 2020, 17, 398-405.	0.9	9
26	Origin and cross-species transmission of bat coronaviruses in China. <i>Nature Communications</i> , 2020, 11, 4235.	5.8	264
27	Lancet COVID-19 Commission Statement on the occasion of the 75th session of the UN General Assembly. <i>Lancet</i> , The, 2020, 396, 1102-1124.	6.3	117
28	No Evidence of Coronaviruses or Other Potentially Zoonotic Viruses in Sunda pangolins (Manis) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 38	0.9	53
29	Nipah virus dynamics in bats and implications for spillover to humans. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 29190-29201.	3.3	119
30	Pandemic COVID-19 Joins History's Pandemic Legion. <i>MBio</i> , 2020, 11, .	1.8	100
31	Sustainable development must account for pandemic risk. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 3888-3892.	3.3	223
32	A qualitative study of zoonotic risk factors among rural communities in southern China. <i>International Health</i> , 2020, 12, 77-85.	0.8	27
33	Escaping Pandora's Box " Another Novel Coronavirus. <i>New England Journal of Medicine</i> , 2020, 382, 1293-1295.	13.9	203
34	Changing Contact Patterns Over Disease Progression: Nipah Virus as a Case Study. <i>Journal of Infectious Diseases</i> , 2020, 222, 438-442.	1.9	4
35	Statement in support of the scientists, public health professionals, and medical professionals of China combatting COVID-19. <i>Lancet</i> , The, 2020, 395, e42-e43.	6.3	182
36	United States wildlife and wildlife product imports from 2000-2014. <i>Scientific Data</i> , 2020, 7, 22.	2.4	33

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37	A strategy to prevent future epidemics similar to the 2019-nCoV outbreak. <i>Biosafety and Health</i> , 2020, 2, 6-8.	1.2	102
38	Make science evolve into a One Health approach to improve health and security: a white paper. <i>One Health Outlook</i> , 2020, 2, 6.	1.4	42
39	Failing Efforts to Mitigate Climate Change are a Futile Band-Aid that will not Stop Other Elephants Filling the Room. <i>EcoHealth</i> , 2020, 17, 421-423.	0.9	2
40	Incorporating Health Outcomes into Land-Use Planning. <i>EcoHealth</i> , 2019, 16, 627-637.	0.9	7
41	Transmission of Nipah Virus – 14 Years of Investigations in Bangladesh. <i>New England Journal of Medicine</i> , 2019, 380, 1804-1814.	13.9	114
42	Using healthcare-seeking behaviour to estimate the number of Nipah outbreaks missed by hospital-based surveillance in Bangladesh. <i>International Journal of Epidemiology</i> , 2019, 48, 1219-1227.	0.9	21
43	Human-animal interactions and bat coronavirus spillover potential among rural residents in Southern China. <i>Biosafety and Health</i> , 2019, 1, 84-90.	1.2	94
44	Isolation and Full-Genome Characterization of Nipah Viruses from Bats, Bangladesh. <i>Emerging Infectious Diseases</i> , 2019, 25, 166-170.	2.0	32
45	Climate Change and Health: Transcending Silos to Find Solutions. <i>Annals of Global Health</i> , 2018, 81, 445.	0.8	32
46	The Global Virome Project. <i>Science</i> , 2018, 359, 872-874.	6.0	324
47	Serological Evidence of Bat SARS-Related Coronavirus Infection in Humans, China. <i>Virologica Sinica</i> , 2018, 33, 104-107.	1.2	219
48	Fatal swine acute diarrhoea syndrome caused by an HKU2-related coronavirus of bat origin. <i>Nature</i> , 2018, 556, 255-258.	13.7	565
49	Characterization of the Spatial and Temporal Distribution of Nipah Virus Spillover Events in Bangladesh, 2007–2013. <i>Journal of Infectious Diseases</i> , 2018, 217, 1390-1394.	1.9	20
50	The Economics of Infectious Disease, Trade and Pandemic Risk. <i>EcoHealth</i> , 2018, 15, 241-243.	0.9	15
51	Comparative analysis of rodent and small mammal viromes to better understand the wildlife origin of emerging infectious diseases. <i>Microbiome</i> , 2018, 6, 178.	4.9	150
52	Serologic and behavioral risk survey of workers with wildlife contact in China. <i>PLoS ONE</i> , 2018, 13, e0194647.	1.1	8
53	Spotted Fever Rickettsiosis in a Wildlife Researcher in Sabah, Malaysia: A Case Study. <i>Tropical Medicine and Infectious Disease</i> , 2018, 3, 29.	0.9	4
54	Nipah Virus Contamination of Hospital Surfaces during Outbreaks, Bangladesh, 2013–2014. <i>Emerging Infectious Diseases</i> , 2018, 24, 15-21.	2.0	39

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55	Middle East Respiratory Syndrome Coronavirus Antibodies in Dromedary Camels, Bangladesh, 2015. <i>Emerging Infectious Diseases</i> , 2018, 24, 926-928.	2.0	19
56	The Economic Case for a Pandemic Fund. <i>EcoHealth</i> , 2018, 15, 244-258.	0.9	28
57	Building a global atlas of zoonotic viruses. <i>Bulletin of the World Health Organization</i> , 2018, 96, 292-294.	1.5	42
58	Hotspots of canine leptospirosis in the United States of America. <i>Veterinary Journal</i> , 2017, 222, 29-35.	0.6	36
59	Does the impact of biodiversity differ between emerging and endemic pathogens? The need to separate the concepts of hazard and risk. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2017, 372, 20160129.	1.8	58
60	Host and viral traits predict zoonotic spillover from mammals. <i>Nature</i> , 2017, 546, 646-650.	13.7	811
61	One Health, emerging infectious diseases and wildlife: two decades of progress?. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2017, 372, 20160167.	1.8	334
62	Potential Sympatric Vectors and Mammalian Hosts of Venezuelan Equine Encephalitis Virus in Southern Mexico. <i>Journal of Wildlife Diseases</i> , 2017, 53, 657.	0.3	16
63	Global hotspots and correlates of emerging zoonotic diseases. <i>Nature Communications</i> , 2017, 8, 1124.	5.8	645
64	Corrigendum to "Global correlates of emerging zoonoses: Anthropogenic, environmental, and biodiversity risk factors" [Int. J. Infect. Dis. 53 (Supplement) (December 2016) 21]. <i>International Journal of Infectious Diseases</i> , 2017, 58, 68.	1.5	0
65	A guide for ecologists: Detecting the role of disease in faunal declines and managing population recovery. <i>Biological Conservation</i> , 2017, 214, 136-146.	1.9	33
66	Economic growth, urbanization, globalization, and the risks of emerging infectious diseases in China: A review. <i>Ambio</i> , 2017, 46, 18-29.	2.8	183
67	Evaluating one health: Are we demonstrating effectiveness?. <i>One Health</i> , 2017, 3, 5-10.	1.5	79
68	Global patterns in coronavirus diversity. <i>Virus Evolution</i> , 2017, 3, vex012.	2.2	310
69	Genetically Diverse Filoviruses in <i>Rousettus</i> and <i>Eonycteris</i> spp. Bats, China, 2009 and 2015. <i>Emerging Infectious Diseases</i> , 2017, 23, 482-486.	2.0	64
70	Convergence of Humans, Bats, Trees, and Culture in Nipah Virus Transmission, Bangladesh. <i>Emerging Infectious Diseases</i> , 2017, 23, 1446-1453.	2.0	76
71	Discovery of a rich gene pool of bat SARS-related coronaviruses provides new insights into the origin of SARS coronavirus. <i>PLoS Pathogens</i> , 2017, 13, e1006698.	2.1	797
72	Cross-sectional surveillance of Middle East respiratory syndrome coronavirus (MERS-CoV) in dromedary camels and other mammals in Egypt, August 2015 to January 2016. <i>Eurosurveillance</i> , 2017, 22,	3.9	41

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73	Genetically Diverse Filoviruses in Rousettus and Eonycteris spp. Bats, China, 2009 and 2015. Emerging Infectious Diseases, 2017, 23, 482-486.	2.0	1
74	Nipah Virus Transmission from Bats to Humans Associated with Drinking Traditional Liquor Made from Date Palm Sap, Bangladesh, 2011-2014. Emerging Infectious Diseases, 2016, 22, 664-670.	2.0	104
75	Evolving epidemiology of Nipah virus infection in Bangladesh: evidence from outbreaks during 2010-2011. Epidemiology and Infection, 2016, 144, 371-380.	1.0	42
76	Bat Severe Acute Respiratory Syndrome-Like Coronavirus WIV1 Encodes an Extra Accessory Protein, ORFX, Involved in Modulation of the Host Immune Response. Journal of Virology, 2016, 90, 6573-6582.	1.5	57
77	Avoiding catastrophes: seeking synergies among the public health, environmental protection, and human security sectors. The Lancet Global Health, 2016, 4, e680-e681.	2.9	7
78	Viral Diversity, Prey Preference, and Bartonella Prevalence in Desmodus rotundus in Guatemala. EcoHealth, 2016, 13, 761-774.	0.9	37
79	Our Collective Vomit. EcoHealth, 2016, 13, 604-605.	0.9	0
80	Quantifying Global Drivers of Zoonotic Bat Viruses: A Process-Based Perspective. American Naturalist, 2016, 187, E53-E64.	1.0	56
81	Investigating Rare Risk Factors for Nipah Virus in Bangladesh: 2001-2012. EcoHealth, 2016, 13, 720-728.	0.9	41
82	Future Earth Health Knowledge-Action Network. Public Health Reviews, 2016, 37, 25.	1.3	4
83	A Last Waltz for Burke, Wills, and King. EcoHealth, 2016, 13, 821-823.	0.9	0
84	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.	0.9	10
85	Fugong virus, a novel hantavirus harbored by the small oriental vole (Eothenomys eleusis) in China. Virology Journal, 2016, 13, 27.	1.4	16
86	Reply to "Complexities of Estimating Evolutionary Rates in Viruses". Journal of Virology, 2016, 90, 2156-2156.	1.5	0
87	Isolation and Characterization of a Novel Bat Coronavirus Closely Related to the Direct Progenitor of Severe Acute Respiratory Syndrome Coronavirus. Journal of Virology, 2016, 90, 3253-3256.	1.5	221
88	Emerging Viral Zoonoses from Wildlife Associated with Animal-Based Food Systems: Risks and Opportunities. , 2016, , 31-57.		11
89	The Emerging Amphibian Fungal Disease, Chytridiomycosis: A Key Example of the Global Phenomenon of Wildlife Emerging Infectious Diseases. Microbiology Spectrum, 2016, 4, .	1.2	18
90	Integrated cluster- and case-based surveillance for detecting stage III zoonotic pathogens: an example of Nipah virus surveillance in Bangladesh. Epidemiology and Infection, 2015, 143, 1922-1930.	1.0	21

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91	Future Earth and EcoHealth: A New Paradigm Toward Global Sustainability and Health. <i>EcoHealth</i> , 2015, 12, 553-554.	0.9	9
92	Spillover and pandemic properties of zoonotic viruses with high host plasticity. <i>Scientific Reports</i> , 2015, 5, 14830.	1.6	238
93	Molecular evidence of Ebola Reston virus infection in Philippine bats. <i>Virology Journal</i> , 2015, 12, 107.	1.4	71
94	Global Avian Influenza Surveillance in Wild Birds: A Strategy to Capture Viral Diversity. <i>Emerging Infectious Diseases</i> , 2015, 21, e1-7.	2.0	46
95	Macacine Herpesvirus 1 in Long-Tailed Macaques, Malaysia, 2009–2011. <i>Emerging Infectious Diseases</i> , 2015, 21, 1107-1113.	2.0	19
96	Exposure-Based Screening for Nipah Virus Encephalitis, Bangladesh. <i>Emerging Infectious Diseases</i> , 2015, 21, 349-351.	2.0	13
97	<i>Bartonella</i> spp. in a Puerto Rican Bat Community. <i>Journal of Wildlife Diseases</i> , 2015, 51, 274-278.	0.3	26
98	Diversity of coronavirus in bats from Eastern Thailand. <i>Virology Journal</i> , 2015, 12, 57.	1.4	70
99	Joint China-US Call for Employing a Transdisciplinary Approach to Emerging Infectious Diseases. <i>EcoHealth</i> , 2015, 12, 555-559.	0.9	3
100	Beyond Ebola: lessons to mitigate future pandemics. <i>The Lancet Global Health</i> , 2015, 3, e354-e355.	2.9	42
101	Reservoir Host Immune Responses to Emerging Zoonotic Viruses. <i>Cell</i> , 2015, 160, 20-35.	13.5	114
102	Serological Evidence of <i>Coxiella burnetii</i> Infection in Cattle and Goats in Bangladesh. <i>EcoHealth</i> , 2015, 12, 354-358.	0.9	11
103	Metacommunity and phylogenetic structure determine wildlife and zoonotic infectious disease patterns in time and space. <i>Ecology and Evolution</i> , 2015, 5, 865-873.	0.8	64
104	Non-random patterns in viral diversity. <i>Nature Communications</i> , 2015, 6, 8147.	5.8	65
105	Integrating invasion and disease in the risk assessment of live bird trade. <i>Diversity and Distributions</i> , 2015, 21, 101-110.	1.9	17
106	Global biogeography of human infectious diseases. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 12746-12751.	3.3	109
107	Evolutionary Dynamics and Global Diversity of Influenza A Virus. <i>Journal of Virology</i> , 2015, 89, 10993-11001.	1.5	46
108	Targeting Transmission Pathways for Emerging Zoonotic Disease Surveillance and Control. <i>Vector-Borne and Zoonotic Diseases</i> , 2015, 15, 432-437.	0.6	119

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109	Middle East Respiratory Syndrome Coronavirus Infection in Dromedary Camels in Saudi Arabia. MBio, 2014, 5, e00884-14.	1.8	359
110	Economic optimization of a global strategy to address the pandemic threat. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 18519-18523.	3.3	113
111	Evidence for Retrovirus and Paramyxovirus Infection of Multiple Bat Species in China. Viruses, 2014, 6, 2138-2154.	1.5	25
112	Serological Evidence of Henipavirus Exposure in Cattle, Goats and Pigs in Bangladesh. PLoS Neglected Tropical Diseases, 2014, 8, e3302.	1.3	57
113	Reply to "Concerns About Misinterpretation of Recent Scientific Data Implicating Dromedary Camels in Epidemiology of Middle East Respiratory Syndrome (MERS)". MBio, 2014, 5, e01482-14.	1.8	4
114	Middle East Respiratory Syndrome Coronavirus Quasispecies That Include Homologues of Human Isolates Revealed through Whole-Genome Analysis and Virus Cultured from Dromedary Camels in Saudi Arabia. MBio, 2014, 5, e01146-14.	1.8	140
115	Evidence for henipavirus spillover into human populations in Africa. Nature Communications, 2014, 5, 5342.	5.8	143
116	Coccidian parasites of red squirrels (<i>Sciurus vulgaris</i>) and grey squirrels (<i>Sciurus</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 462 Td	0.2	5
117	Merging Economics and Epidemiology to Improve the Prediction and Management of Infectious Disease. EcoHealth, 2014, 11, 464-475.	0.9	87
118	Ebola Economics: The Case for an Upstream Approach to Disease Emergence. EcoHealth, 2014, 11, 451-452.	0.9	8
119	Prevalence and Diversity of Avian Haematozoan Parasites in Wetlands of Bangladesh. Journal of Parasitology Research, 2014, 2014, 1-12.	0.5	19
120	Roosting behaviour and habitat selection of <i>Pteropus giganteus</i> reveal potential links to Nipah virus epidemiology. Journal of Applied Ecology, 2014, 51, 376-387.	1.9	58
121	Giardia, HIV, and Nature's Horrifying Beauty. EcoHealth, 2014, 11, 277-278.	0.9	0
122	Detection of diverse novel astroviruses from small mammals in China. Journal of General Virology, 2014, 95, 2442-2449.	1.3	33
123	We Are All Animals. EcoHealth, 2014, 11, 145-146.	0.9	0
124	Dengue Virus in Bats from Southeastern Mexico. American Journal of Tropical Medicine and Hygiene, 2014, 91, 129-131.	0.6	40
125	The Role of Landscape Composition and Configuration on <i>Pteropus giganteus</i> Roosting Ecology and Nipah Virus Spillover Risk in Bangladesh. American Journal of Tropical Medicine and Hygiene, 2014, 90, 247-255.	0.6	62
126	Bird migration and avian influenza: A comparison of hydrogen stable isotopes and satellite tracking methods. Ecological Indicators, 2014, 45, 266-273.	2.6	25

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127	Characterization of recombinant H9N2 influenza viruses isolated from wild ducks in China. <i>Veterinary Microbiology</i> , 2013, 166, 327-336.	0.8	16
128	Predicted and observed mortality from vector-borne disease in wildlife: West Nile virus and small songbirds. <i>Biological Conservation</i> , 2013, 165, 79-85.	1.9	25
129	Isolation and characterization of a bat SARS-like coronavirus that uses the ACE2 receptor. <i>Nature</i> , 2013, 503, 535-538.	13.7	1,439
130	Pathogens, Pests, and Economics: Drivers of Honey Bee Colony Declines and Losses. <i>EcoHealth</i> , 2013, 10, 434-445.	0.9	187
131	Interdisciplinary approaches to understanding disease emergence: The past, present, and future drivers of Nipah virus emergence. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 3681-3688.	3.3	128
132	Lack of population genetic structure and host specificity in the bat fly, <i>Cyclopodia horsfieldi</i> , across species of <i>Pteropus</i> bats in Southeast Asia. <i>Parasites and Vectors</i> , 2013, 6, 231.	1.0	37
133	Linking the Historical Roots of Environmental Conservation with Human and Wildlife Health. <i>EcoHealth</i> , 2013, 10, 224-227.	0.9	3
134	A Strategy To Estimate Unknown Viral Diversity in Mammals. <i>MBio</i> , 2013, 4, e00598-13.	1.8	320
135	Human ecology in pathogenic landscapes: two hypotheses on how land use change drives viral emergence. <i>Current Opinion in Virology</i> , 2013, 3, 79-83.	2.6	137
136	The Human Environment Interface: Applying Ecosystem Concepts to Health. <i>Current Topics in Microbiology and Immunology</i> , 2013, 365, 83-100.	0.7	9
137	Ebola Virus Antibodies in Fruit Bats, Bangladesh. <i>Emerging Infectious Diseases</i> , 2013, 19, 270-273.	2.0	129
138	Nipah Virus Infection Outbreak with Nosocomial and Corpse-to-Human Transmission, Bangladesh. <i>Emerging Infectious Diseases</i> , 2013, 19, 210-217.	2.0	110
139	Middle East Respiratory Syndrome Coronavirus in Bats, Saudi Arabia. <i>Emerging Infectious Diseases</i> , 2013, 19, 1819-23.	2.0	562
140	Using network theory to identify the causes of disease outbreaks of unknown origin. <i>Journal of the Royal Society Interface</i> , 2013, 10, 20130127.	1.5	1
141	Using network theory to identify the causes of disease outbreaks of unknown origin. <i>Journal of the Royal Society Interface</i> , 2013, 10, 20120904.	1.5	13
142	Predicting Hotspots for Influenza Virus Reassortment. <i>Emerging Infectious Diseases</i> , 2013, 19, 581-588.	2.0	62
143	Bats are a major natural reservoir for hepaciviruses and pegiviruses. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 8194-8199.	3.3	251
144	Ecological Approaches to Studying Zoonoses. <i>Microbiology Spectrum</i> , 2013, 1, .	1.2	3

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145	Risk Factors for Nipah Virus Infection among Pteropid Bats, Peninsular Malaysia. <i>Emerging Infectious Diseases</i> , 2013, 19, 51-60.	2.0	44
146	Targeting Surveillance for Zoonotic Virus Discovery. <i>Emerging Infectious Diseases</i> , 2013, 19, 743-747.	2.0	37
147	Quantifying Trends in Disease Impact to Produce a Consistent and Reproducible Definition of an Emerging Infectious Disease. <i>PLoS ONE</i> , 2013, 8, e69951.	1.1	19
148	Foraging Behaviour and Landscape Utilisation by the Endangered Golden-Crowned Flying Fox (<i>Acerodon jubatus</i>), The Philippines. <i>PLoS ONE</i> , 2013, 8, e79665.	1.1	15
149	The Human Environment Interface: Applying Ecosystem Concepts to Health. <i>Current Topics in Microbiology and Immunology</i> , 2013, , 83-100.	0.7	6
150	Duration of Maternal Antibodies against Canine Distemper Virus and Hendra Virus in Pteropid Bats. <i>PLoS ONE</i> , 2013, 8, e67584.	1.1	37
151	Identification of a Novel Cetacean Polyomavirus from a Common Dolphin (<i>Delphinus delphis</i>) with Tracheobronchitis. <i>PLoS ONE</i> , 2013, 8, e68239.	1.1	18
152	Metapopulation Dynamics Enable Persistence of Influenza A, Including A/H5N1, in Poultry. <i>PLoS ONE</i> , 2013, 8, e80091.	1.1	13
153	Preventing Pandemics Via International Development: A Systems Approach. <i>PLoS Medicine</i> , 2012, 9, e1001354.	3.9	37
154	Agricultural intensification, priming for persistence and the emergence of Nipah virus: a lethal bat-borne zoonosis. <i>Journal of the Royal Society Interface</i> , 2012, 9, 89-101.	1.5	245
155	Emergence of Fatal Avian Influenza in New England Harbor Seals. <i>MBio</i> , 2012, 3, e00166-12.	1.8	161
156	Anatomy of a pandemic. <i>Lancet</i> , The, 2012, 380, 1883-1884.	6.3	28
157	Prediction and prevention of the next pandemic zoonosis. <i>Lancet</i> , The, 2012, 380, 1956-1965.	6.3	744
158	Mainstreaming One Health. <i>EcoHealth</i> , 2012, 9, 107-110.	0.9	79
159	Corn, Flour, Blue. <i>EcoHealth</i> , 2012, 9, 234-236.	0.9	0
160	Two Views of the New China. <i>EcoHealth</i> , 2012, 9, 367-369.	0.9	0
161	A New Species of <i>Eimeria</i> (Apicomplexa: Eimeriidae) from the Endangered Pink Pigeon, <i>Nesoenas mayeri</i> (Prévost, 1843) Cheke, 2005 (Columbiformes) in Mauritius. <i>African Zoology</i> , 2012, 47, 369-372.	0.2	5
162	Towards a Better Integration of Global Health and Biodiversity in the New Sustainable Development Goals Beyond Rio+20. <i>EcoHealth</i> , 2012, 9, 381-385.	0.9	27

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163	Novel, panzootic and hybrid genotypes of amphibian chytridiomycosis associated with the bullfrog trade. <i>Molecular Ecology</i> , 2012, 21, 5162-5177.	2.0	227
164	Biodiversity and ecosystem services science for a sustainable planet: the DIVERSITAS vision for 2012-2020. <i>Current Opinion in Environmental Sustainability</i> , 2012, 4, 101-105.	3.1	62
165	We Are All But Wardian Cases. <i>EcoHealth</i> , 2012, 9, 504-505.	0.9	0
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