

Impacts of biodiversity on the emergence and transmis

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Replication of bacteriophage G13 DNA in dna mutants of Escherichia coli. Nucleic Acids and Protein Synthesis, 1978, 520, 505-511.	1.7	1
2	A Community-Ecology Framework for Understanding Vector and Vector-Borne Disease Dynamics. Israel Journal of Ecology and Evolution, 2010, 56, 251-262.	0.2	8
3	An Insect Nidovirus Emerging from a Primary Tropical Rainforest. MBio, 2011, 2, e00077-11.	1.8	100
4	Tropical amphibian populations experience higher disease risk in natural habitats. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 9893-9898.	3.3	144
5	A dilution effect in the emerging amphibian pathogen <i>Batrachochytrium dendrobatidis</i> . Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 16322-16326.	3.3	98
6	Food security: why is biodiversity important?. International Forestry Review, 2011, 13, 265-274.	0.3	125
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9	Pathogen impacts on plant communities: unifying theory, concepts, and empirical work. Ecological Monographs, 2011, 81, 429-441.	2.4	224
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18	Horizon scan of global conservation issues for 2011. Trends in Ecology and Evolution, 2011, 26, 10-16.	4.2	213
19	Endangered species and a threatened discipline: behavioural ecology. Trends in Ecology and Evolution, 2011, 26, 111-118.	4.2	78

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21	Prevalence of Anaplasma phagocytophilum in Ixodes ricinus in Bavarian public parks, Germany. Ticks and Tick-borne Diseases, 2011, 2, 196-203.	1.1	51
22	Infectious Diseases, Biodiversity and Global Changes: How the Biodiversity Sciences May Help. , 0, , .		3
23	Guidance on the environmental risk assessment of plant pests. EFSA Journal, 2011, 9, 2460.	0.9	34
24	The Mechanism of Attributing Economic Value to the Natural Environment. SSRN Electronic Journal, 2011, , .	0.4	0
25	Global Impact of Mosquito Biodiversity, Human Vector-Borne Diseases and Environmental Change. , 0, , .		23
26	Biodiversity of Borrelia burgdorferi Strains in Tissues of Lyme Disease Patients. PLoS ONE, 2011, 6, e22926.	1.1	64
27	Fine-Scale Variation in Vector Host Use and Force of Infection Drive Localized Patterns of West Nile Virus Transmission. PLoS ONE, 2011, 6, e23767.	1.1	106
28	Invasion and Persistence of Infectious Agents in Fragmented Host Populations. PLoS ONE, 2011, 6, e24006.	1.1	11
29	Establishment of Normal Gut Microbiota Is Compromised under Excessive Hygiene Conditions. PLoS ONE, 2011, 6, e28284.	1.1	120
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37	A back-arc in time. Nature, 2011, 469, 170-171.	13.7	0

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47	Improved Knockout Methodology Reveals That Frog Virus 3 Mutants Lacking either the 18K Immediate-Early Gene or the Truncated vIF-2 Gene Are Defective for Replication and Growth <i>In Vivo</i> . <i>Journal of Virology</i> , 2011, 85, 11131-11138.	1.5	44
48	Reservoir Competence of Various Rodents for the Lyme Disease Spirochete <i>Borrelia spielmanii</i> . <i>Applied and Environmental Microbiology</i> , 2011, 77, 3565-3570.	1.4	43
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54	Effect of Biodiversity Changes in Disease Risk: Exploring Disease Emergence in a Plant-Virus System. <i>PLoS Pathogens</i> , 2012, 8, e1002796.	2.1	105
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57	Leptospirosis in American Samoa – Estimating and Mapping Risk Using Environmental Data. <i>PLoS Neglected Tropical Diseases</i> , 2012, 6, e1669.	1.3	58
58	Host Life History Strategy, Species Diversity, and Habitat Influence <i>Trypanosoma cruzi</i> Vector Infection in Changing Landscapes. <i>PLoS Neglected Tropical Diseases</i> , 2012, 6, e1884.	1.3	100
59	Climate change and animal diseases: making the case for adaptation. <i>Animal Health Research Reviews</i> , 2012, 13, 209-222.	1.4	5
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62	Leptospirosis in American Samoa 2010: Epidemiology, Environmental Drivers, and the Management of Emergence. <i>American Journal of Tropical Medicine and Hygiene</i> , 2012, 86, 309-319.	0.6	52
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73	The Global Burden of Human Parasites: Who and Where are They? How are They Transmitted?. <i>Journal of Parasitology</i> , 2012, 98, 1056-1064.	0.3	33
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76	Prediction and prevention of the next pandemic zoonosis. <i>Lancet, The</i> , 2012, 380, 1956-1965.	6.3	744
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94	Unraveling plant–animal diversity relationships: a meta-regression analysis. <i>Ecology</i> , 2012, 93, 2115-2124.	1.5	114
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130	Relationship between pace of life and immune responses in wild rodents. <i>Oikos</i> , 2012, 121, 1483-1492.	1.2	114

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131	Deciphering arboviral emergence within insular ecosystems. <i>Infection, Genetics and Evolution</i> , 2012, 12, 1333-1339.	1.0	31
132	A curve of thresholds governs plague epizootics in Central Asia. <i>Ecology Letters</i> , 2012, 15, 554-560.	3.0	32
133	Emergence of new leptospiral serovars in American Samoa - ascertainment or ecological change?. <i>BMC Infectious Diseases</i> , 2012, 12, 19.	1.3	25
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140	Flow cytometry, microscopy, and DNA analysis as complementary phytoplankton screening methods in ballast water treatment studies. <i>Journal of Applied Phycology</i> , 2013, 25, 1047-1053.	1.5	17
141	The biodiversity hypothesis and allergic disease: world allergy organization position statement. <i>World Allergy Organization Journal</i> , 2013, 6, 3.	1.6	282
142	Experimental Evidence for American Bullfrog (<i>Lithobates catesbeianus</i>) Susceptibility to Chytrid Fungus (<i>Batrachochytrium dendrobatidis</i>). <i>EcoHealth</i> , 2013, 10, 166-171.	0.9	44
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145	Mitigating amphibian chytridiomycosis with bioaugmentation: characteristics of effective probiotics and strategies for their selection and use. <i>Ecology Letters</i> , 2013, 16, 807-820.	3.0	239
146	Wildlife disease ecology in changing landscapes: Mesopredator release and toxoplasmosis. <i>International Journal for Parasitology: Parasites and Wildlife</i> , 2013, 2, 110-118.	0.6	62
147	Climate, vegetation, introduced hosts and trade shape a global wildlife pandemic. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2013, 280, 20122506.	1.2	99
148	Does high biodiversity reduce the risk of Lyme disease invasion?. <i>Parasites and Vectors</i> , 2013, 6, 195.	1.0	40

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156	The Impact of Community Organization on Vector-Borne Pathogens. <i>American Naturalist</i> , 2013, 181, 1-11.	1.0	115
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158	<i>Echinostoma trivolvis</i> (Digenea: Echinostomatidae) second intermediate host preference matches host suitability. <i>Parasitology Research</i> , 2013, 112, 799-805.	0.6	10
159	Biodiversity decreases disease through predictable changes in host community competence. <i>Nature</i> , 2013, 494, 230-233.	13.7	288
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161	Reinstating soil microbial diversity in agroecosystems: The need of the hour for sustainability and health. <i>Agriculture, Ecosystems and Environment</i> , 2013, 164, 181-182.	2.5	44
162	Experimental viral evolution reveals major histocompatibility complex polymorphisms as the primary host factors controlling pathogen adaptation and virulence. <i>Genes and Immunity</i> , 2013, 14, 365-372.	2.2	17
163	It's a myth that protection against disease is a strong and general service of biodiversity conservation: Response to Ostfeld and Keesing. <i>Trends in Ecology and Evolution</i> , 2013, 28, 503-504.	4.2	46
164	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
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166	Does habitat disturbance increase infectious disease risk for primates?. <i>Ecology Letters</i> , 2013, 16, 656-663.	3.0	78
167	Factors driving hantavirus emergence in Europe. <i>Current Opinion in Virology</i> , 2013, 3, 92-99.	2.6	64

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168	The Relevance of Forest Fragmentation on the Incidence of Human Babesiosis: Investigating the Landscape Epidemiology of an Emerging Tick-Borne Disease. <i>Vector-Borne and Zoonotic Diseases</i> , 2013, 13, 250-255.	0.6	16
169	A meta-analysis suggesting that the relationship between biodiversity and risk of zoonotic pathogen transmission is idiosyncratic. <i>Ecology Letters</i> , 2013, 16, 679-686.	3.0	211
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