

Kate E Langwig

List of Publications by Year in descending order

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Version: 2024-02-01

30
papers

2,953
citations

257429

24
h-index

454934

30
g-index

34
all docs

34
docs citations

34
times ranked

2183
citing authors

#	ARTICLE	IF	CITATIONS
1	An Emerging Disease Causes Regional Population Collapse of a Common North American Bat Species. <i>Science</i> , 2010, 329, 679-682.	12.6	735
2	Sociality, density dependence and microclimates determine the persistence of populations suffering from a novel fungal disease, white-nose syndrome. <i>Ecology Letters</i> , 2012, 15, 1050-1057.	6.4	299
3	Disease alters macroecological patterns of North American bats. <i>Global Ecology and Biogeography</i> , 2015, 24, 741-749.	5.8	206
4	Host and pathogen ecology drive the seasonal dynamics of a fungal disease, white-nose syndrome. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2015, 282, 20142335.	2.6	181
5	Context-dependent conservation responses to emerging wildlife diseases. <i>Frontiers in Ecology and the Environment</i> , 2015, 13, 195-202.	4.0	147
6	Bacteria Isolated from Bats Inhibit the Growth of <i>Pseudogymnoascus destructans</i> , the Causative Agent of White-Nose Syndrome. <i>PLoS ONE</i> , 2015, 10, e0121329.	2.5	120
7	Ecology and impacts of white-nose syndrome on bats. <i>Nature Reviews Microbiology</i> , 2021, 19, 196-210.	28.6	107
8	Pathogen dynamics during invasion and establishment of white-nose syndrome explain mechanisms of host persistence. <i>Ecology</i> , 2017, 98, 624-631.	3.2	100
9	Drivers of variation in species impacts for a multi-host fungal disease of bats. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2016, 371, 20150456.	4.0	92
10	Invasion Dynamics of White-Nose Syndrome Fungus, Midwestern United States, 2012–2014. <i>Emerging Infectious Diseases</i> , 2015, 21, 1023-1026.	4.3	88
11	Resistance in persisting bat populations after white-nose syndrome invasion. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2017, 372, 20160044.	4.0	86
12	Deconstructing the Bat Skin Microbiome: Influences of the Host and the Environment. <i>Frontiers in Microbiology</i> , 2016, 7, 1753.	3.5	81
13	Long-Term Persistence of <i>Pseudogymnoascus destructans</i> , the Causative Agent of White-Nose Syndrome, in the Absence of Bats. <i>EcoHealth</i> , 2015, 12, 330-333.	2.0	68
14	Risk factors associated with mortality from white-nose syndrome among hibernating bat colonies. <i>Biology Letters</i> , 2011, 7, 950-953.	2.3	62
15	Little Brown Myotis Persist Despite Exposure to White-Nose Syndrome. <i>Journal of Fish and Wildlife Management</i> , 2011, 2, 190-195.	0.9	62
16	Efficacy of a probiotic bacterium to treat bats affected by the disease white-nose syndrome. <i>Journal of Applied Ecology</i> , 2017, 54, 701-708.	4.0	59
17	Widespread Bat White-Nose Syndrome Fungus, Northeastern China. <i>Emerging Infectious Diseases</i> , 2015, 22, 140-142.	4.3	54
18	Cryptic connections illuminate pathogen transmission within community networks. <i>Nature</i> , 2018, 563, 710-713.	27.8	54

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19	Environmental reservoir dynamics predict global infection patterns and population impacts for the fungal disease white-nose syndrome. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 7255-7262.	7.1	53
20	Field trial of a probiotic bacteria to protect bats from white-nose syndrome. <i>Scientific Reports</i> , 2019, 9, 9158.	3.3	50
21	Moving Beyond Too Little, Too Late: Managing Emerging Infectious Diseases in Wild Populations Requires International Policy and Partnerships. <i>EcoHealth</i> , 2015, 12, 404-407.	2.0	45
22	Host persistence or extinction from emerging infectious disease: insights from white-nose syndrome in endemic and invading regions. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2016, 283, 20152861.	2.6	40
23	Vaccine Effects on Heterogeneity in Susceptibility and Implications for Population Health Management. <i>MBio</i> , 2017, 8, .	4.1	32
24	Integral Projection Models for host-parasite systems with an application to amphibian chytrid fungus. <i>Methods in Ecology and Evolution</i> , 2016, 7, 1182-1194.	5.2	28
25	Continued preference for suboptimal habitat reduces bat survival with white-nose syndrome. <i>Nature Communications</i> , 2021, 12, 166.	12.8	19
26	Limited available evidence supports theoretical predictions of reduced vaccine efficacy at higher exposure dose. <i>Scientific Reports</i> , 2019, 9, 3203.	3.3	18
27	On the Fly: Interactions Between Birds, Mosquitoes, and Environment That Have Molded West Nile Virus Genomic Structure Over Two Decades. <i>Journal of Medical Entomology</i> , 2019, 56, 1467-1474.	1.8	17
28	Host traits and environment interact to determine persistence of bat populations impacted by white-nose syndrome. <i>Ecology Letters</i> , 2022, 25, 483-497.	6.4	15
29	Impact of censusing and research on wildlife populations. <i>Conservation Science and Practice</i> , 2020, 2, e264.	2.0	10
30	Mobility and infectiousness in the spatial spread of an emerging fungal pathogen. <i>Journal of Animal Ecology</i> , 2021, 90, 1134-1141.	2.8	10