

Kim M Pepin

List of Publications by Year in descending order

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Version: 2024-02-01

73
papers

2,317
citations

331670

21
h-index

254184

43
g-index

79
all docs

79
docs citations

79
times ranked

3028
citing authors

#	ARTICLE	IF	CITATIONS
1	Influence of biotic and abiotic factors on home range size and shape of invasive wild pigs (<i>Sus</i> Tj ETQq1 1 0.784314 rgBT/Overlo	3.4	10
2	Accounting for animal movement improves vaccination strategies against wildlife disease in heterogeneous landscapes. <i>Ecological Applications</i> , 2022, 32, e2568.	3.8	10
3	A model for leveraging animal movement to understand spatio-temporal disease dynamics. <i>Ecology Letters</i> , 2022, 25, 1290-1304.	6.4	16
4	An efficient method of evaluating multiple concurrent management actions on invasive populations. <i>Ecological Applications</i> , 2022, 32, e2623.	3.8	8
5	Optimizing management of invasions in an uncertain world using dynamic spatial models. <i>Ecological Applications</i> , 2022, 32, e2628.	3.8	5
6	Adaptive risk-based targeted surveillance for foreign animal diseases at the wildlife-livestock interface. <i>Transboundary and Emerging Diseases</i> , 2022, 69, .	3.0	6
7	Transmission of antibiotic resistance at the wildlife-livestock interface. <i>Communications Biology</i> , 2022, 5, .	4.4	17
8	Defining an epidemiological landscape that connects movement ecology to pathogen transmission and pace-of-life. <i>Ecology Letters</i> , 2022, 25, 1760-1782.	6.4	18
9	Multi-level movement response of invasive wild pigs (<i>Sus scrofa</i>) to removal. <i>Pest Management Science</i> , 2021, 77, 85-95.	3.4	9
10	Embracing Dynamic Models for Gene Drive Management. <i>Trends in Biotechnology</i> , 2021, 39, 211-214.	9.3	10
11	Effects of social structure and management on risk of disease establishment in wild pigs. <i>Journal of Animal Ecology</i> , 2021, 90, 820-833.	2.8	21
12	Social structure defines spatial transmission of African swine fever in wild boar. <i>Journal of the Royal Society Interface</i> , 2021, 18, 20200761.	3.4	14
13	A framework for surveillance of emerging pathogens at the human-animal interface: Pigs and coronaviruses as a case study. <i>Preventive Veterinary Medicine</i> , 2021, 188, 105281.	1.9	8
14	Behavioral state resource selection in invasive wild pigs in the Southeastern United States. <i>Scientific Reports</i> , 2021, 11, 6924.	3.3	23
15	Strategic testing approaches for targeted disease monitoring can be used to inform pandemic decision-making. <i>PLoS Biology</i> , 2021, 19, e3001307.	5.6	9
16	Spatial variation in direct and indirect contact rates at the wildlife-livestock interface for informing disease management. <i>Preventive Veterinary Medicine</i> , 2021, 194, 105423.	1.9	13
17	Quantifying Transmission Between Wild and Domestic Populations. <i>Wildlife Research Monographs</i> , 2021, , 369-409.	0.9	1
18	How do genetic relatedness and spatial proximity shape African swine fever infections in wild boar?. <i>Transboundary and Emerging Diseases</i> , 2021, , .	3.0	5

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19	Predicting functional responses in agro-ecosystems from animal movement data to improve management of invasive pests. <i>Ecological Applications</i> , 2020, 30, e02015.	3.8	14
20	Inferring seasonal infection risk at population and regional scales from serology samples. <i>Ecology</i> , 2020, 101, e02882.	3.2	6
21	Factors Affecting Bait Site Visitation: Area of Influence of Baits. <i>Wildlife Society Bulletin</i> , 2020, 44, 362-371.	1.6	8
22	Optimal spatial prioritization of control resources for elimination of invasive species under demographic uncertainty. <i>Ecological Applications</i> , 2020, 30, e02126.	3.8	14
23	A Rapid Population Assessment Method for Wild Pigs Using Baited Cameras at 3 Study Sites. <i>Wildlife Society Bulletin</i> , 2020, 44, 372-382.	1.6	6
24	Ecological drivers of African swine fever virus persistence in wild boar populations: Insight for control. <i>Ecology and Evolution</i> , 2020, 10, 2846-2859.	1.9	60
25	A comparison of cost and quality of three methods for estimating density for wild pig (<i>Sus scrofa</i>). <i>Scientific Reports</i> , 2020, 10, 2047.	3.3	11
26	Variation in host home range size decreases rabies vaccination effectiveness by increasing the spatial spread of rabies virus. <i>Journal of Animal Ecology</i> , 2020, 89, 1375-1386.	2.8	28
27	Optimal bait density for delivery of acute toxicants to vertebrate pests. <i>Journal of Pest Science</i> , 2020, 93, 723-735.	3.7	7
28	Linking mosquito surveillance to dengue fever through Bayesian mechanistic modeling. <i>PLoS Neglected Tropical Diseases</i> , 2020, 14, e0008868.	3.0	8
29	Epidemic growth rates and host movement patterns shape management performance for pathogen spillover at the wildlife-livestock interface. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2019, 374, 20180343.	4.0	10
30	Improving risk assessment of the emergence of novel influenza A viruses by incorporating environmental surveillance. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2019, 374, 20180346.	4.0	11
31	Confronting models with data: the challenges of estimating disease spillover. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2019, 374, 20180435.	4.0	30
32	Not all surveillance data are created equal—A multi-method dynamic occupancy approach to determine rabies elimination from wildlife. <i>Journal of Applied Ecology</i> , 2019, 56, 2551-2561.	4.0	19
33	Individual-Level Antibody Dynamics Reveal Potential Drivers of Influenza A Seasonality in Wild Pig Populations. <i>Integrative and Comparative Biology</i> , 2019, 59, 1231-1242.	2.0	8
34	Rabies Surveillance Identifies Potential Risk Corridors and Enables Management Evaluation. <i>Viruses</i> , 2019, 11, 1006.	3.3	15
35	Ecological interventions to prevent and manage zoonotic pathogen spillover. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2019, 374, 20180342.	4.0	102
36	BOARD INVITED REVIEW: Prospects for improving management of animal disease introductions using disease-dynamic models. <i>Journal of Animal Science</i> , 2019, 97, 2291-2307.	0.5	17

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37	Modelling multi-species and multi-mode contact networks: Implications for persistence of bovine tuberculosis at the wildlife-livestock interface. <i>Journal of Applied Ecology</i> , 2019, 56, 1471-1481.	4.0	24
38	Accounting for heterogeneous invasion rates reveals management impacts on the spatial expansion of an invasive species. <i>Ecosphere</i> , 2019, 10, e02657.	2.2	18
39	Exposure of a population of invasive wild pigs to simulated toxic bait containing biomarker: implications for population reduction. <i>Pest Management Science</i> , 2019, 75, 1140-1149.	3.4	15
40	Quantifying site-level usage and certainty of absence for an invasive species through occupancy analysis of camera-trap data. <i>Biological Invasions</i> , 2018, 20, 877-890.	2.4	14
41	Accounting for observation processes across multiple levels of uncertainty improves inference of species distributions and guides adaptive sampling of environmental DNA. <i>Ecology and Evolution</i> , 2018, 8, 10879-10892.	1.9	25
42	Costs and effectiveness of damage management of an overabundant species (<i>Sus scrofa</i>) using aerial gunning. <i>Wildlife Research</i> , 2018, 45, 696.	1.4	20
43	Inferring infection hazard in wildlife populations by linking data across individual and population scales. <i>Ecology Letters</i> , 2017, 20, 275-292.	6.4	50
44	Estimating population density for disease risk assessment: The importance of understanding the area of influence of traps using wild pigs as an example. <i>Preventive Veterinary Medicine</i> , 2017, 141, 33-37.	1.9	18
45	Quantifying drivers of wild pig movement across multiple spatial and temporal scales. <i>Movement Ecology</i> , 2017, 5, 14.	2.8	75
46	Comment on: "Blood does not buy goodwill: allowing culling increases poaching of a large carnivore". <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2017, 284, 20161459.	2.6	12
47	Effects of scale of movement, detection probability, and true population density on common methods of estimating population density. <i>Scientific Reports</i> , 2017, 7, 9446.	3.3	47
48	Efficiency of different spatial and temporal strategies for reducing vertebrate pest populations. <i>Ecological Modelling</i> , 2017, 365, 106-118.	2.5	13
49	The persistence of multiple strains of avian influenza in live bird markets. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2017, 284, 20170715.	2.6	11
50	Potential effects of incorporating fertility control into typical culling regimes in wild pig populations. <i>PLoS ONE</i> , 2017, 12, e0183441.	2.5	33
51	Predicting spatial spread of rabies in skunk populations using surveillance data reported by the public. <i>PLoS Neglected Tropical Diseases</i> , 2017, 11, e0005822.	3.0	17
52	Inferring invasive species abundance using removal data from management actions. <i>Ecological Applications</i> , 2016, 26, 2339-2346.	3.8	36
53	Contact heterogeneities in feral swine: implications for disease management and future research. <i>Ecosphere</i> , 2016, 7, e01230.	2.2	35
54	Evaluating wildlife-cattle contact rates to improve the understanding of dynamics of bovine tuberculosis transmission in Michigan, USA. <i>Preventive Veterinary Medicine</i> , 2016, 135, 28-36.	1.9	19

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55	Disease-emergence dynamics and control in a socially-structured wildlife species. <i>Scientific Reports</i> , 2016, 6, 25150.	3.3	16
56	Invasion of two tick-borne diseases across New England: harnessing human surveillance data to capture underlying ecological invasion processes. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2016, 283, 20160834.	2.6	26
57	Utility of mosquito surveillance data for spatial prioritization of vector control against dengue viruses in three Brazilian cities. <i>Parasites and Vectors</i> , 2015, 8, 98.	2.5	18
58	Deer response to exclusion from stored cattle feed in Michigan, USA. <i>Preventive Veterinary Medicine</i> , 2015, 121, 159-164.	1.9	16
59	Improving pandemic influenza risk assessment. <i>ELife</i> , 2014, 3, e03883.	6.0	53
60	Using quantitative disease dynamics as a tool for guiding response to avian influenza in poultry in the United States of America. <i>Preventive Veterinary Medicine</i> , 2014, 113, 376-397.	1.9	19
61	Fitting outbreak models to data from many small norovirus outbreaks. <i>Epidemics</i> , 2014, 6, 18-29.	3.0	21
62	Multiannual patterns of influenza A transmission in Chinese live bird market systems. <i>Influenza and Other Respiratory Viruses</i> , 2013, 7, 97-107.	3.4	41
63	Persistence of black-tailed prairie-dog populations affected by plague in northern Colorado, USA. <i>Ecology</i> , 2013, 94, 1572-1583.	3.2	23
64	Cost-effectiveness of Novel System of Mosquito Surveillance and Control, Brazil. <i>Emerging Infectious Diseases</i> , 2013, 19, 542-550.	4.3	46
65	Anticipating the Prevalence of Avian Influenza Subtypes H9 and H5 in Live-Bird Markets. <i>PLoS ONE</i> , 2013, 8, e56157.	2.5	10
66	Geographic Variation in the Relationship between Human Lyme Disease Incidence and Density of Infected Host-Seeking <i>Ixodes scapularis</i> Nymphs in the Eastern United States. <i>American Journal of Tropical Medicine and Hygiene</i> , 2012, 86, 1062-1071.	1.4	141
67	Identifying genetic markers of adaptation for surveillance of viral host jumps. <i>Nature Reviews Microbiology</i> , 2010, 8, 802-813.	28.6	138
68	Synthesizing within-host and population-level selective pressures on viral populations: the impact of adaptive immunity on viral immune escape. <i>Journal of the Royal Society Interface</i> , 2010, 7, 1311-1318.	3.4	23
69	Epidemic Dynamics at the Human-Animal Interface. <i>Science</i> , 2009, 326, 1362-1367.	12.6	554
70	Genomic evolution in a virus under specific selection for host recognition. <i>Infection, Genetics and Evolution</i> , 2008, 8, 825-834.	2.3	47
71	Density-Dependent Competitive Suppression of Sylvatic Dengue Virus by Endemic Dengue Virus in Cultured Mosquito Cells. <i>Vector-Borne and Zoonotic Diseases</i> , 2008, 8, 821-828.	1.5	21
72	Variable Pleiotropic Effects From Mutations at the Same Locus Hamper Prediction of Fitness From a Fitness Component. <i>Genetics</i> , 2006, 172, 2047-2056.	2.9	29

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73	Molecular Cloning of Horse Hsp90 cDNA and Its Comparative Analysis with Other Vertebrate Hsp90 Sequences.. Journal of Veterinary Medical Science, 2001, 63, 115-124.	0.9	13