Meggan E Craft

List of Publications by Year in descending order

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172207 168136 3,509 92 29 53 citations h-index g-index papers 102 102 102 4512 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Parasites as conservation tools. Conservation Biology, 2022, 36, .	2.4	24
2	Hunting alters viral transmission and evolution in a large carnivore. Nature Ecology and Evolution, 2022, 6, 174-182.	3.4	5
3	How to study parasites and host migration: a roadmap for empiricists. Biological Reviews, 2022, 97, 1161-1178.	4.7	6
4	Social associations in common carp ($\langle i \rangle$ Cyprinus carpio $\langle i \rangle$): Insights from induced feeding aggregations for targeted management strategies. Ecology and Evolution, 2022, 12, e8666.	0.8	3
5	Paradoxes and synergies: Optimizing management of a deadly virus in an endangered carnivore. Journal of Applied Ecology, 2022, 59, 1548-1558.	1.9	3
6	Asymmetric host movement reshapes local disease dynamics in metapopulations. Scientific Reports, 2022, 12, .	1.6	0
7	Defining an epidemiological landscape that connects movement ecology to pathogen transmission and paceâ€ofâ€ife. Ecology Letters, 2022, 25, 1760-1782.	3.0	18
8	Tradeâ€offs with telemetryâ€derived contact networks for infectious disease studies in wildlife. Methods in Ecology and Evolution, 2021, 12, 76-87.	2.2	26
9	Bidirectional interactions between host social behaviour and parasites arise through ecological and evolutionary processes. Parasitology, 2021, 148, 274-288.	0.7	30
10	Group density, disease, and season shape territory size and overlap of social carnivores. Journal of Animal Ecology, 2021, 90, 87-101.	1.3	12
11	Parasite intensity and the evolution of migratory behavior. Ecology, 2021, 102, e03229.	1.5	8
12	Host relatedness and landscape connectivity shape pathogen spread in the puma, a large secretive carnivore. Communications Biology, 2021, 4, 12.	2.0	20
13	Patterns and processes of pathogen exposure in gray wolves across North America. Scientific Reports, 2021, 11, 3722.	1.6	6
14	Importance of anthropogenic sources at shaping the antimicrobial resistance profile of a peri-urban mesocarnivore. Science of the Total Environment, 2021, 764, 144166.	3.9	9
15	Network connectivity of Minnesota waterbodies and implications for aquatic invasive species prevention. Biological Invasions, 2021, 23, 3231-3242.	1.2	11
16	Comparison of Antimicrobial-Resistant Escherichia coli Isolates from Urban Raccoons and Domestic Dogs. Applied and Environmental Microbiology, 2021, 87, e0048421.	1.4	6
17	Infection risk varies within urbanized landscapes: the case of coyotes and heartworm. Parasites and Vectors, 2021, 14, 464.	1.0	3
18	Lessons from movement ecology for the return to work: Modeling contacts and the spread of COVID-19. PLoS ONE, 2021, 16, e0242955.	1.1	6

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19	Microbial associations and spatial proximity predict North American moose (<i>Alces alces</i>) gastrointestinal community composition. Journal of Animal Ecology, 2020, 89, 817-828.	1.3	16
20	A mechanistic, stigmergy model of territory formation in solitary animals: Territorial behavior can dampen disease prevalenceÂbut increase persistence. PLoS Computational Biology, 2020, 16, e1007457.	1.5	9
21	Characterization of antimicrobial resistance genes in Enterobacteriaceae carried by suburban mesocarnivores and locally owned and stray dogs. Zoonoses and Public Health, 2020, 67, 460-466.	0.9	7
22	Does the virus cross the road? Viral phylogeographic patterns among bobcat populations reflect a history of urban development. Evolutionary Applications, 2020, 13, 1806-1817.	1.5	7
23	Crossâ€species transmission and evolutionary dynamics of canine distemper virus during a spillover in African lions of Serengeti National Park. Molecular Ecology, 2020, 29, 4308-4321.	2.0	18
24	Using host traits to predict reservoir host species of rabies virus. PLoS Neglected Tropical Diseases, 2020, 14, e0008940.	1.3	29
25	Genotype and sex-based host variation in behaviour and susceptibility drives population disease dynamics. Proceedings of the Royal Society B: Biological Sciences, 2020, 287, 20201653.	1.2	4
26	Title is missing!. , 2020, 16, e1007457.		0
27	Title is missing!. , 2020, 16, e1007457.		0
28	Title is missing!., 2020, 16, e1007457.		0
29	Title is missing!. , 2020, 16, e1007457.		0
30	Title is missing!., 2020, 16, e1007457.		0
31	Title is missing!. , 2020, 16, e1007457.		O
32	How to make more from exposure data? An integrated machine learning pipeline to predict pathogen exposure. Journal of Animal Ecology, 2019, 88, 1447-1461.	1.3	33
33	Mapping parasite transmission risk from white-tailed deer to a declining moose population. European Journal of Wildlife Research, 2019, 65, 1.	0.7	13
34	City sicker? A metaâ€analysis of wildlife health and urbanization. Frontiers in Ecology and the Environment, 2019, 17, 575-583.	1.9	114
35	Feline immunodeficiency virus in puma: Estimation of force of infection reveals insights into transmission. Ecology and Evolution, 2019, 9, 11010-11024.	0.8	7
36	Urbanization impacts apex predator gene flow but not genetic diversity across an urbanâ€rural divide. Molecular Ecology, 2019, 28, 4926-4940.	2.0	23

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37	Host migration strategy is shaped by forms of parasite transmission and infection cost. Journal of Animal Ecology, 2019, 88, 1601-1612.	1.3	16
38	Emerging human infectious diseases and the links to global food production. Nature Sustainability, 2019, 2, 445-456.	11.5	362
39	Endemic infection can shape exposure to novel pathogens: Pathogen coâ€occurrence networks in the Serengeti lions. Ecology Letters, 2019, 22, 904-913.	3.0	14
40	Transmission ecology of canine parvovirus in a multi-host, multi-pathogen system. Proceedings of the Royal Society B: Biological Sciences, 2019, 286, 20182772.	1.2	26
41	Speciesâ€specific spatiotemporal patterns of leopard, lion and tiger attacks on humans. Journal of Applied Ecology, 2019, 56, 585-593.	1.9	24
42	Characterization of swine movements in the United States and implications for disease control. Preventive Veterinary Medicine, 2019, 164, 1-9.	0.7	36
43	Model recommendations meet management reality: implementation and evaluation of a network-informed vaccination effort for endangered Hawaiian monk seals. Proceedings of the Royal Society B: Biological Sciences, 2018, 285, 20171899.	1.2	16
44	Managing complexity: Simplifying assumptions of foot-and-mouth disease models for swine. Transboundary and Emerging Diseases, 2018, 65, 1307-1317.	1.3	17
45	Incorporating genomic methods into contact networks to reveal new insights into animal behaviour and infectious disease dynamics. Behaviour, 2018, 155, 759-791.	0.4	16
46	Covariation between the physiological and behavioral components of pathogen transmission: host heterogeneity determines epidemic outcomes. Oikos, 2018, 127, 538-552.	1.2	23
47	Dynamic, spatial models of parasite transmission in wildlife: Their structure, applications and remaining challenges. Journal of Animal Ecology, 2018, 87, 559-580.	1.3	56
48	Towards an ecoâ€phylogenetic framework for infectious disease ecology. Biological Reviews, 2018, 93, 950-970.	4.7	63
49	Challenges and Opportunities Developing Mathematical Models of Shared Pathogens of Domestic and Wild Animals. Veterinary Sciences, 2018, 5, 92.	0.6	14
50	Infectious Disease in Wild Animal Populations: Examining Transmission and Control with Mathematical Models. Advances in Environmental Microbiology, 2018, , 239-266.	0.1	1
51	Modeling cost-effectiveness of risk-based bovine tuberculosis surveillance in Minnesota. Preventive Veterinary Medicine, 2018, 159, 1-11.	0.7	15
52	Disease outbreak thresholds emerge from interactions between movement behavior, landscape structure, and epidemiology. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 7374-7379.	3.3	69
53	Pathogens in space: Advancing understanding of pathogen dynamics and disease ecology through landscape genetics. Evolutionary Applications, 2018, 11, 1763-1778.	1.5	37
54	Potential distribution of the viral haemorrhagic septicaemia virus in the Great Lakes region. Journal of Fish Diseases, 2017, 40, 11-28.	0.9	18

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55	Influenza A virus in swine breeding herds: Combination of vaccination and biosecurity practices can reduce likelihood of endemic piglet reservoir. Preventive Veterinary Medicine, 2017, 138, 55-69.	0.7	36
56	Urban landscapes can change virus gene flow and evolution in a fragmentationâ€sensitive carnivore. Molecular Ecology, 2017, 26, 6487-6498.	2.0	40
57	Linking social and spatial networks to viral community phylogenetics reveals subtypeâ€specific transmission dynamics in African lions. Journal of Animal Ecology, 2017, 86, 1469-1482.	1.3	22
58	Using contact networks to explore mechanisms of parasite transmission in wildlife. Biological Reviews, 2017, 92, 389-409.	4.7	136
59	Interactions between domestic and wild carnivores around the greater Serengeti ecosystem. Animal Conservation, 2017, 20, 193-204.	1.5	17
60	Seasonality and pathogen transmission in pastoral cattle contact networks. Royal Society Open Science, 2017, 4, 170808.	1.1	44
61	Inferring the Ecological Niche of Toxoplasma gondii and Bartonella spp. in Wild Felids. Frontiers in Veterinary Science, 2017, 4, 172.	0.9	3
62	Advances and Limitations of Disease Biogeography Using Ecological Niche Modeling. Frontiers in Microbiology, 2016, 07, 1174.	1.5	105
63	"One Health―or Three? Publication Silos Among the One Health Disciplines. PLoS Biology, 2016, 14, e1002448.	2.6	84
64	Parameter Values for Epidemiological Models of Foot-and-Mouth Disease in Swine. Frontiers in Veterinary Science, 2016, 3, 44.	0.9	29
65	Which mechanisms drive seasonal rabies outbreaks in raccoons? A test using dynamic social network models. Journal of Applied Ecology, 2016, 53, 804-813.	1.9	34
66	Is pathogen exposure spatially autocorrelated? Patterns of pathogens in puma (Puma concolor) and bobcat (Lynx rufus). Ecosphere, 2016, 7, e01558.	1.0	12
67	Declining Prevalence of Disease Vectors Under Climate Change. Scientific Reports, 2016, 6, 39150.	1.6	46
68	Host behaviour–parasite feedback: an essential link between animal behaviour and disease ecology. Proceedings of the Royal Society B: Biological Sciences, 2016, 283, 20153078.	1.2	112
69	Evaluating empirical contact networks as potential transmission pathways for infectious diseases. Journal of the Royal Society Interface, 2016, 13, 20160166.	1.5	41
70	Mosquitoes in Moose Country: A Mosquito Survey of Northern Minnesota. Journal of the American Mosquito Control Association, 2016, 32, 83-90.	0.2	2
71	Network analysis of cattle movements in Uruguay: Quantifying heterogeneity for risk-based disease surveillance and control. Preventive Veterinary Medicine, 2016, 123, 12-22.	0.7	58
72	Raccoon contact networks predict seasonal susceptibility to rabies outbreaks and limitations of vaccination. Journal of Animal Ecology, 2015, 84, 1720-1731.	1.3	67

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73	From network analysis to risk analysisâ€"An approach to risk-based surveillance for bovine tuberculosis in Minnesota, US. Preventive Veterinary Medicine, 2015, 118, 328-340.	0.7	19
74	Dynamics of a morbillivirus at the domestic–wildlife interface: Canine distemper virus in domestic dogs and lions. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 1464-1469.	3.3	128
75	Infectious disease transmission and contact networks in wildlife and livestock. Philosophical Transactions of the Royal Society B: Biological Sciences, 2015, 370, 20140107.	1.8	251
76	The sociality–health–fitness nexus: synthesis, conclusions and future directions. Philosophical Transactions of the Royal Society B: Biological Sciences, 2015, 370, 20140115.	1.8	41
77	Episodic outbreaks of small mammals influence predator community dynamics in an east African savanna ecosystem. Oikos, 2014, 123, 1014-1024.	1.2	28
78	Risk-Based Management of Viral Hemorrhagic Septicemia Virus in Minnesota. North American Journal of Fisheries Management, 2014, 34, 373-379.	0.5	9
79	Mathematical Modeling of Influenza A Virus Dynamics within Swine Farms and the Effects of Vaccination. PLoS ONE, 2014, 9, e106177.	1.1	48
80	Canine Distemper Virus (CDV) in Another Big Cat: Should CDV Be Renamed Carnivore Distemper Virus?. MBio, 2013, 4, e00702-13.	1.8	46
81	Epidemiological effects of group size variation in social species. Journal of the Royal Society Interface, 2013, 10, 20130206.	1.5	22
82	Asynchronous food-web pathways could buffer the response of Serengeti predators to El Niño Southern Oscillation. Ecology, 2013, 94, 1123-1130.	1.5	27
83	Estimating the Probability of a Major Outbreak from the Timing of Early Cases: An Indeterminate Problem?. PLoS ONE, 2013, 8, e57878.	1.1	21
84	FIV diversity: FIVPle subtype composition may influence disease outcome in African lions. Veterinary Immunology and Immunopathology, 2011, 143, 338-346.	0.5	27
85	Longâ€ŧerm trends in carnivore abundance using distance sampling in Serengeti National Park, Tanzania. Journal of Applied Ecology, 2011, 48, 1490-1500.	1.9	65
86	Disease transmission in territorial populations: the small-world network of Serengeti lions. Journal of the Royal Society Interface, 2011, 8, 776-786.	1.5	121
87	Network Models: An Underutilized Tool in Wildlife Epidemiology?. Interdisciplinary Perspectives on Infectious Diseases, 2011, 2011, 1-12.	0.6	107
88	Does size matter? An investigation of habitat use across a carnivore assemblage in the Serengeti, Tanzania. Journal of Animal Ecology, 2010, 79, 1012-1022.	1.3	27
89	Distinguishing epidemic waves from disease spillover in a wildlife population. Proceedings of the Royal Society B: Biological Sciences, 2009, 276, 1777-1785.	1.2	80
90	Capture and rapid handling of jackals (Canis mesomelas and Canis adustus) without chemical immobilization. African Journal of Ecology, 2008, 46, 214-216.	0.4	3

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91	Dynamics of a multihost pathogen in a carnivore community. Journal of Animal Ecology, 2008, 77, 1257-1264.	1.3	79
92	Exploring reservoir dynamics: a case study of rabies in the Serengeti ecosystem. Journal of Applied Ecology, 2008, 45, 1246-1257.	1.9	166