

Roderick B Gagne

List of Publications by Year in descending order

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Version: 2024-02-01

37
papers

523
citations

686830

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752256

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42
all docs

42
docs citations

42
times ranked

719
citing authors

#	ARTICLE	IF	CITATIONS
1	SARS-CoV-2 evolution in animals suggests mechanisms for rapid variant selection. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	3.3	69
2	Climate change and conservation of endemic amphidromous fishes in Hawaiian streams. Endangered Species Research, 2012, 16, 261-272.	1.2	43
3	Genetic source-sink dynamics among naturally structured and anthropogenically fragmented puma populations. Conservation Genetics, 2019, 20, 215-227.	0.8	33
4	Overcoming urban stream syndrome: Trophic flexibility confers resilience in a Hawaiian stream fish. Freshwater Biology, 2018, 63, 492-502.	1.2	25
5	Urbanization reduces genetic connectivity in bobcats (<i>Lynx rufus</i>) at both intra- and interpopulation spatial scales. Molecular Ecology, 2019, 28, 5068-5085.	2.0	24
6	Parasites as conservation tools. Conservation Biology, 2022, 36, .	2.4	24
7	Molecular and Morphological Evidence of Distinct Evolutionary Lineages of <i>Awaous guamensis</i> in Hawai'i and Guam. Copeia, 2012, 2012, 293-300.	1.4	23
8	Urbanization impacts apex predator gene flow but not genetic diversity across an urban-rural divide. Molecular Ecology, 2019, 28, 4926-4940.	2.0	23
9	Microsatellite records for volume 8, issue 1. Conservation Genetics Resources, 2016, 8, 43-81.	0.4	22
10	Measures of effective population size in sea otters reveal special considerations for wide-ranging species. Evolutionary Applications, 2018, 11, 1779-1790.	1.5	20
11	Host relatedness and landscape connectivity shape pathogen spread in the puma, a large secretive carnivore. Communications Biology, 2021, 4, 12.	2.0	20
12	Spread of an introduced parasite across the Hawaiian archipelago independent of its introduced host. Freshwater Biology, 2015, 60, 311-322.	1.2	18
13	Frequent cross-species transmissions of foamy virus between domestic and wild felids. Virus Evolution, 2020, 6, vez058.	2.2	17
14	Variation in Intra-individual Lentiviral Evolution Rates: a Systematic Review of Human, Nonhuman Primate, and Felid Species. Journal of Virology, 2019, 93, .	1.5	15
15	MrIML: Multi-response interpretable machine learning to model genomic landscapes. Molecular Ecology Resources, 2021, 21, 2766-2781.	2.2	12
16	Colonization and demographic expansion of freshwater fauna across the Hawaiian archipelago. Journal of Evolutionary Biology, 2016, 29, 2054-2069.	0.8	11
17	Mutual dilution of infection by an introduced parasite in native and invasive stream fishes across Hawaii. Parasitology, 2016, 143, 1605-1614.	0.7	9
18	Optimal DNA extractions from blood on preservation paper limits conservation genomic but not conservation genetic applications. Journal for Nature Conservation, 2018, 46, 89-96.	0.8	9

#	ARTICLE	IF	CITATIONS
19	The Expectations and Challenges of Wildlife Disease Research in the Era of Genomics: Forecasting with a Horizon Scan-like Exercise. <i>Journal of Heredity</i> , 2019, 110, 261-274.	1.0	9
20	Pronghorn population genomics show connectivity in the core of their range. <i>Journal of Mammalogy</i> , 2020, 101, 1061-1071.	0.6	9
21	Invasion of the Hawaiian Islands by a parasite infecting imperiled stream fishes. <i>Ecography</i> , 2018, 41, 528-539.	2.1	8
22	Phylogeography of the widespread creek chub <i>Semotilus atromaculatus</i> (Cypriniformes: Leuciscidae). <i>Journal of Fish Biology</i> , 2018, 93, 778-791.	0.7	8
23	Mitogenomes and relatedness do not predict frequency of tool-use by sea otters. <i>Biology Letters</i> , 2017, 13, 20160880.	1.0	7
24	Bighorn Sheep Genetic Structure in Wyoming Reflects Geography and Management. <i>Journal of Wildlife Management</i> , 2020, 84, 1072-1090.	0.7	7
25	Functional connectivity in a continuously distributed, migratory species as revealed by landscape genomics. <i>Ecography</i> , 2021, 44, 987.	2.1	7
26	Population genomic diversity and structure at the discontinuous southern range of the Great Gray Owl in North America. <i>Conservation Genetics</i> , 2020, 21, 693-706.	0.8	6
27	Hunting alters viral transmission and evolution in a large carnivore. <i>Nature Ecology and Evolution</i> , 2022, 6, 174-182.	3.4	5
28	Multi-population puma connectivity could restore genomic diversity to at-risk coastal populations in California. <i>Evolutionary Applications</i> , 2022, 15, 286-299.	1.5	5
29	Parasitism of a native Hawaiian stream fish by an introduced nematode increases with declining precipitation across a natural rainfall gradient. <i>Ecology of Freshwater Fish</i> , 2016, 25, 476-486.	0.7	4
30	Novel hybrid finds a peri-urban niche: Allen's Hummingbirds in southern California. <i>Conservation Genetics</i> , 2020, 21, 989-998.	0.8	4
31	Altered lentiviral infection dynamics follow genetic rescue of the Florida panther. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2019, 286, 20191689.	1.2	3
32	Translocations maintain genetic diversity and increase connectivity in sea otters, <i>Enhydra lutris</i> . <i>Marine Mammal Science</i> , 2021, 37, 1475-1497.	0.9	3
33	Genetic Characterization of <i>Microsporium canis</i> Clinical Isolates in the United States. <i>Journal of Fungi (Basel, Switzerland)</i> , 2022, 8, 676.	1.5	3
34	A natural laboratory to elucidate the evolution of endogenous-exogenous retroviral interactions. <i>Molecular Ecology</i> , 2021, 30, 2473-2476.	2.0	0
35	Viral Sequences Recovered From Puma Tooth DNA Reconstruct Statewide Viral Phylogenies. <i>Frontiers in Ecology and Evolution</i> , 2021, 9, .	1.1	0
36	INVESTIGATING ASSOCIATIONS AMONG RELATEDNESS, GENETIC DIVERSITY, AND CAUSES OF MORTALITY IN SOUTHERN SEA OTTERS (<i>ENHYDRA LUTRIS NEREIS</i>). <i>Journal of Wildlife Diseases</i> , 2022, 58, .	0.3	0

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37	Modelling the role of predation on disease burdens of prey. <i>Journal of Animal Ecology</i> , 2022, 91, 1330-1333.	1.3	0