Nusha Keyghobadi

List of Publications by Year
in descending order

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47 papers	2,289 citations	430754 18 h-index	223716 46 g-index
47 all docs	47 docs citations	47 times ranked	2800 citing authors

#	Article	IF	Citations
1	High genetic drift in endangered northern peripheral populations of the Behr's hairstreak butterfly (<scp>Satyrium behrii</scp>). Insect Conservation and Diversity, 2021, 14, 403-411.	1.4	3
2	Evolutionary history of the Persian squirrel (Sciurus anomalus): It emerged on the Eurasian continent in the Miocene. Zoologischer Anzeiger, 2020, 287, 17-24.	0.4	6
3	Demographic fluctuations lead to rapid and cyclic shifts in genetic structure among populations of an alpine butterfly, Parnassius smintheus. Journal of Evolutionary Biology, 2020, 33, 668-681.	0.8	9
4	Direct estimates of metapopulation capacity from dispersal show high interannual variability, but little effect of recent forest encroachment on network persistence. Landscape Ecology, 2020, 35, 675-688.	1.9	3
5	The permanent inhabitant of the oak trees: phylogeography and genetic structure of the Persian squirrel (Sciurus anomalus). Biological Journal of the Linnean Society, 2019, 127, 197-212.	0.7	19
6	Dispersing male Parnassius smintheus butterflies are more strongly affected by forest matrix than are females. Insect Science, 2019, 26, 932-944.	1.5	6
7	Bed bugs: The move to humans as hosts. Facets, 2019, 4, 105-110.	1.1	1
8	Host association influences variation at salivary protein genes in the bat ectoparasite Cimex adjunctus. Journal of Evolutionary Biology, 2018, 31, 753-763.	0.8	4
9	Limited genetic evidence for host plantâ€related differentiation in the Western cherry fruit fly, Rhagoletis indifferens. Entomologia Experimentalis Et Applicata, 2018, 166, 739-751.	0.7	5
10	Analysis of genetic diversity in a peatland specialist butterfly suggests an important role for habitat quality and small habitat patches. Conservation Genetics, 2018, 19, 1109-1121.	0.8	3
11	Flight morphology corresponds to both surrounding landscape structure and local patch conditions in a highly specialized peatland butterfly (Lycaena epixanthe). Ecological Entomology, 2018, 43, 629-639.	1.1	8
12	Comparative analysis of landscape effects on spatial genetic structure of the big brown bat and one of its cimicid ectoparasites. Ecology and Evolution, 2017, 7, 8210-8219.	0.8	9
13	Population structure in two geographically sympatric and congeneric ectoparasites (Cimex adjunctus) Tj ETQq1 1 0.784314 rgBT /Ov 95, 901-907.	0.4	6
14	Host association and selection on salivary protein genes in bed bugs and related blood-feeding ectoparasites. Royal Society Open Science, 2017, 4, 170446.	1.1	7
15	Connectivity rescues genetic diversity after a demographic bottleneck in a butterfly population network. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 10914-10919.	3.3	67
16	Range-wide genetic structure and demographic history in the bat ectoparasite Cimex adjunctus. BMC Evolutionary Biology, 2016, 16, 268.	3.2	23
17	Swallowtail butterflies show positive edge responses predicted by resource use. Landscape Ecology, 2016, 31, 2115-2131.	1.9	5
18	Landscape genetics in a changing world: disentangling historical and contemporary influences and inferring change. Molecular Ecology, 2015, 24, 6021-6040.	2.0	210

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19	The relative influence of habitat amount and configuration on genetic structure across multiple spatial scales. Ecology and Evolution, 2015, 5, 73-86.	0.8	14
20	Landscape structure and the genetic effects of a population collapse. Proceedings of the Royal Society B: Biological Sciences, 2014, 281, 20141798.	1.2	17
21	Population genetic structure of the western cherry fruit flyRhagoletis indifferens(Diptera:) Tj ETQq1 1 0.784314 rgBT /Overlock	0.7	9
22	Ten years of abundance data within a spatial population network of the alpine butterfly,Parnassius smintheus. Ecology, 2014, 95, 2985.	1.5	5
23	Historical specimens reveal past relationships and current conservation status of populations in a declining species: the regal fritillary butterfly. Insect Conservation and Diversity, 2013, 6, 234-242.	1.4	15
24	Mating success and oviposition of a butterfly are not affected by non-lethal tissue sampling. Journal of Insect Conservation, 2013, 17, 859-864.	0.8	9
25	Invasion genetics of <scp>A</scp>merican cherry fruit fly in <scp>E</scp>urope and signals of hybridization with the <scp>E</scp>uropean cherry fruit fly. Entomologia Experimentalis Et Applicata, 2013, 147, 61-72.	0.7	20
26	A call for more transparent reporting of error rates: the quality of AFLP data in ecological and evolutionary research. Molecular Ecology, 2012, 21, 5911-5917.	2.0	32
27	The Pitcher Plant Flesh Fly Exhibits a Mixture of Patchy and Metapopulation Attributes. Journal of Heredity, 2012, 103, 703-710.	1.0	6
28	From broadscale patterns to fineâ€scale processes: habitat structure influences genetic differentiation in the pitcher plant midge across multiple spatial scales. Molecular Ecology, 2012, 21, 223-236.	2.0	15
29	Fine-scale genetic structure of an endangered population of the Mormon metalmark butterfly (Apodemia mormo) revealed using AFLPs. Conservation Genetics, 2011, 12, 991-1001.	0.8	19
30	Effects of different methods of non-lethal tissue sampling on butterflies. Ecological Entomology, 2011, 36, 301-308.	1.1	32
31	Successful analysis of AFLPs from non-lethally sampled wing tissues in butterflies. Conservation Genetics, 2009, 10, 2021-2024.	0.8	12
32	Characterization of microsatellite loci for the western cherry fruit fly, Rhagoletis indifferens (Diptera: Tephritidae). Molecular Ecology Resources, 2009, 9, 1025-1028.	2.2	18
33	Characterization of microsatellite loci for the pitcher plant midge, Metriocnemus knabi Coq. (Diptera: Chironomidae). Molecular Ecology Resources, 2009, 9, 1388-1391.	2.2	5
34	The genetic implications of habitat fragmentation for animalsThis review is one of a series dealing with some aspects of the impact of habitat fragmentation on animals and plants. This series is one of several virtual symposia focussing on ecological topics that will be published in the Journal from time to time.. Canadian Journal of Zoology, 2007, 85, 1049-1064.	0.4	333
35	Fine-scale population genetic structure of a wildlife disease vector: the southern house mosquito on the island of Hawaii. Molecular Ecology, 2006, 15, 3919-3930.	2.0	27
36	Remnant populations of the regal fritillary (Speyeria idalia) in Pennsylvania: Local genetic structure in a high gene flow species. Conservation Genetics, 2006, 7, 309-313.	0.8	17

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37	Cross-species comparison of microsatellite loci in the Culex pipiens complex and beyond. Molecular Ecology Notes, 2005, 5, 697-700.	1.7	53
38	Genetic differentiation and gene flow among populations of the alpine butterfly, Parnassius smintheus, vary with landscape connectivity. Molecular Ecology, 2005, 14, 1897-1909.	2.0	115
39	Among- and within-patch components of genetic diversity respond at different rates to habitat fragmentation: an empirical demonstration. Proceedings of the Royal Society B: Biological Sciences, 2005, 272, 553-560.	1.2	121
40	Emerging Vectors in the Culex pipiens Complex. Science, 2004, 303, 1535-1538.	6.0	438
41	Microsatellite loci from the northern house mosquito (Culex pipiens), a principal vector of West Nile virus in North America. Molecular Ecology Notes, 2003, 4, 20-22.	1.7	47
42	The Effects of Isolation, Habitat Area and Resources on the Abundance, Density and Movement of the Butterfly Parnassius smintheus. American Midland Naturalist, 2003, 150, 26-36.	0.2	33
43	Isolation of novel microsatellite loci in the Rocky Mountain apollo butterfly, Parnassius smintheus. Hereditas, 2002, 136, 247-250.	0.5	22
44	ALPINEPARNASSIUSBUTTERFLY DISPERSAL: EFFECTS OF LANDSCAPE AND POPULATION SIZE. Ecology, 2000, 81, 1642-1653.	1.5	191
45	Alpine Parnassius Butterfly Dispersal: Effects of Landscape and Population Size. Ecology, 2000, 81, 1642.	1.5	85
46	Influence of landscape on the population genetic structure of the alpine butterfly Parnassius smintheus (Papilionidae). Molecular Ecology, 1999, 8, 1481-1495.	2.0	185
47	Microsatellites for the at-risk Mottled Duskywing butterfly, Erynnis martialis. Conservation Genetics Resources, 0, , .	0.4	0