

# Adalgisa Caccone

## List of Publications by Year in descending order

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123  
papers

4,747  
citations

87843

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128225

60  
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127  
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127  
docs citations

127  
times ranked

5319  
citing authors

#	ARTICLE	IF	CITATIONS
1	HUMAN IMPACTS HAVE SHAPED HISTORICAL AND RECENT EVOLUTION IN <i>Aedes aegypti</i> , THE DENGUE AND YELLOW FEVER MOSQUITO. <i>Evolution; International Journal of Organic Evolution</i> , 2014, 68, 514-525.	1.1	225
2	Mitochondrial DNA from Hemlock Woolly Adelgid (Hemiptera: Adelgidae) Suggests Cryptic Speciation and Pinpoints the Source of the Introduction to Eastern North America. <i>Annals of the Entomological Society of America</i> , 2006, 99, 195-203.	1.3	194
3	Colonization and diversification of Galápagos terrestrial fauna: a phylogenetic and biogeographical synthesis. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2008, 363, 3347-3361.	1.8	167
4	PHYLOGEOGRAPHY AND HISTORY OF GIANT GALAPAGOS TORTOISES. <i>Evolution; International Journal of Organic Evolution</i> , 2002, 56, 2052-2066.	1.1	128
5	Global population divergence and admixture of the brown rat ( <i>Rattus norvegicus</i> ). <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2016, 283, 20161762.	1.2	119
6	Independent evolutionary origins of landlocked alewife populations and rapid parallel evolution of phenotypic traits. <i>Molecular Ecology</i> , 2008, 17, 582-597.	2.0	118
7	Multiple Origins of Knockdown Resistance Mutations in the Afrotropical Mosquito Vector <i>Anopheles gambiae</i> . <i>PLoS ONE</i> , 2007, 2, e1243.	1.1	108
8	MOLECULAR BIOGEOGRAPHY OF CAVE LIFE: A STUDY USING MITOCHONDRIAL DNA FROM BATHYSCIINE BEETLES. <i>Evolution; International Journal of Organic Evolution</i> , 2001, 55, 122-130.	1.1	99
9	Analysis of Multiple Tsetse Fly Populations in Uganda Reveals Limited Diversity and Species-Specific Gut Microbiota. <i>Applied and Environmental Microbiology</i> , 2014, 80, 4301-4312.	1.4	95
10	Population genomics of the Asian tiger mosquito, <i>Aedes albopictus</i> : insights into the recent worldwide invasion. <i>Ecology and Evolution</i> , 2017, 7, 10143-10157.	0.8	89
11	Microsatellite analysis of genetic divergence among populations of giant Galápagos tortoises. <i>Molecular Ecology</i> , 2008, 11, 2265-2283.	2.0	88
12	Unravelling the peculiarities of island life: vicariance, dispersal and the diversification of the extinct and extant giant Galápagos tortoises. <i>Molecular Ecology</i> , 2012, 21, 160-173.	2.0	88
13	Phylogeography and Taxonomy of <i>Trypanosoma brucei</i> . <i>PLoS Neglected Tropical Diseases</i> , 2011, 5, e961.	1.3	84
14	Historical DNA analysis reveals living descendants of an extinct species of Galápagos tortoise. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 15464-15469.	3.3	79
15	Giant tortoise genomes provide insights into longevity and age-related disease. <i>Nature Ecology and Evolution</i> , 2019, 3, 87-95.	3.4	79
16	Tracking the return of <i>Aedes aegypti</i> to Brazil, the major vector of the dengue, chikungunya and Zika viruses. <i>PLoS Neglected Tropical Diseases</i> , 2017, 11, e0005653.	1.3	77
17	Title is missing!. <i>Conservation Genetics</i> , 2003, 4, 31-46.	0.8	75
18	Patterns of association between Symbiodinium and members of the <i>Montastraea annularis</i> species complex on spatial scales ranging from within colonies to between geographic regions. <i>Coral Reefs</i> , 2006, 25, 503-512.	0.9	72

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19	Hybridization between a native and introduced predator of Adelgidae: An unintended result of classical biological control. <i>Biological Control</i> , 2012, 63, 359-369.	1.4	72
20	A cryptic taxon of Galápagos tortoise in conservation peril. <i>Biology Letters</i> , 2005, 1, 287-290.	1.0	71
21	Giant tortoises are not so slow: Rapid diversification and biogeographic consensus in the Galapagos. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004, 101, 6514-6519.	3.3	70
22	Genes Record a Prehistoric Volcano Eruption in the Galapagos. <i>Science</i> , 2003, 302, 75-75.	6.0	69
23	Editing nature: Local roots of global governance. <i>Science</i> , 2018, 362, 527-529.	6.0	67
24	Improved reference genome of the arboviral vector <i>Aedes albopictus</i> . <i>Genome Biology</i> , 2020, 21, 215.	3.8	65
25	Ancient and modern colonization of North America by hemlock woolly adelgid, <i>Adelges tsugae</i> (Hemiptera: Adelgidae), an invasive insect from East Asia. <i>Molecular Ecology</i> , 2016, 25, 2065-2080.	2.0	64
26	Lineage fusion in Galápagos giant tortoises. <i>Molecular Ecology</i> , 2014, 23, 5276-5290.	2.0	59
27	Extreme difference in rate of mitochondrial and nuclear DNA evolution in a large ectotherm, Galápagos tortoises. <i>Molecular Phylogenetics and Evolution</i> , 2004, 31, 794-798.	1.2	58
28	Description of a New Galapagos Giant Tortoise Species (Chelonoidis; Testudines: Testudinidae) from Cerro Fatal on Santa Cruz Island. <i>PLoS ONE</i> , 2015, 10, e0138779.	1.1	54
29	Urban population genetics of slum-dwelling rats ( <i>Rattus norvegicus</i> ) in Salvador, Brazil. <i>Molecular Ecology</i> , 2013, 22, 5056-5070.	2.0	52
30	Hybridization masks speciation in the evolutionary history of the Galápagos marine iguana. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2015, 282, 20150425.	1.2	52
31	Genetic analysis of a successful repatriation programme: giant Galápagos tortoises. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2004, 271, 341-345.	1.2	51
32	<i>Trypanosoma brucei gambiense</i> Group 1 Is Distinguished by a Unique Amino Acid Substitution in the HpHb Receptor Implicated in Human Serum Resistance. <i>PLoS Neglected Tropical Diseases</i> , 2012, 6, e1728.	1.3	50
33	Lonesome George is not alone among Galápagos tortoises. <i>Current Biology</i> , 2007, 17, R317-R318.	1.8	49
34	Urban rat races: spatial population genomics of brown rats ( <i>Rattus norvegicus</i> ) compared across multiple cities. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2018, 285, 20180245.	1.2	48
35	<i>Glossina fuscipes</i> populations provide insights for human African trypanosomiasis transmission in Uganda. <i>Trends in Parasitology</i> , 2013, 29, 394-406.	1.5	47
36	Genetic rediscovery of an "extinct" Galápagos giant tortoise species. <i>Current Biology</i> , 2012, 22, R10-R11.	1.8	46

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37	Implications of Microfauna-Host Interactions for Trypanosome Transmission Dynamics in <i>Glossina fuscipes fuscipes</i> in Uganda. <i>Applied and Environmental Microbiology</i> , 2012, 78, 4627-4637.	1.4	45
38	Phylogeography and Population Structure of <i>Glossina fuscipes fuscipes</i> in Uganda: Implications for Control of Tsetse. <i>PLoS Neglected Tropical Diseases</i> , 2010, 4, e636.	1.3	44
39	Using fine-scale spatial genetics of Norway rats to improve control efforts and reduce leptospirosis risk in urban slum environments. <i>Evolutionary Applications</i> , 2017, 10, 323-337.	1.5	43
40	Recovery of a nearly extinct <i>Galapagos</i> tortoise despite minimal genetic variation. <i>Evolutionary Applications</i> , 2013, 6, 377-383.	1.5	42
41	Phylogeographic History and Gene Flow Among Giant <i>Galapagos</i> Tortoises on Southern Isabela Island. <i>Genetics</i> , 2006, 172, 1727-1744.	1.2	40
42	Evolution of <i>kdr</i> haplotypes in worldwide populations of <i>Aedes aegypti</i> : Independent origins of the F1534C <i>kdr</i> mutation. <i>PLoS Neglected Tropical Diseases</i> , 2020, 14, e0008219.	1.3	40
43	The genetic legacy of Lonesome George survives: Giant tortoises with Pinta Island ancestry identified in <i>Galapagos</i> . <i>Biological Conservation</i> , 2013, 157, 225-228.	1.9	39
44	Comparative Genomics Reveals Multiple Genetic Backgrounds of Human Pathogenicity in the <i>Trypanosoma brucei</i> Complex. <i>Genome Biology and Evolution</i> , 2014, 6, 2811-2819.	1.1	39
45	Genomic insights into the ancient spread of Lyme disease across North America. <i>Nature Ecology and Evolution</i> , 2017, 1, 1569-1576.	3.4	39
46	Permanent Genetic Resources added to Molecular Ecology Resources Database 1 December 2010–31 January 2011. <i>Molecular Ecology Resources</i> , 2011, 11, 586-589.	2.2	38
47	Whole genome capture of vector-borne pathogens from mixed DNA samples: a case study of <i>Borrelia burgdorferi</i> . <i>BMC Genomics</i> , 2015, 16, 434.	1.2	38
48	Potential genetic consequences of a recent bottleneck in the Amur tiger of. <i>Conservation Genetics</i> , 2004, 5, 707-713.	0.8	36
49	DNA from the Past Informs Ex Situ Conservation for the Future: An “Extinct” Species of <i>Galapagos</i> Tortoise Identified in Captivity. <i>PLoS ONE</i> , 2010, 5, e8683.	1.1	36
50	High Levels of Genetic Differentiation between Ugandan <i>Glossina fuscipes fuscipes</i> Populations Separated by Lake Kyoga. <i>PLoS Neglected Tropical Diseases</i> , 2008, 2, e242.	1.3	35
51	Genetic Markers of Benzimidazole Resistance among Human Hookworms ( <i>Necator americanus</i> ) in Kintampo North Municipality, Ghana. <i>American Journal of Tropical Medicine and Hygiene</i> , 2019, 100, 351-356.	0.6	35
52	Morphometrics Parallel Genetics in a Newly Discovered and Endangered Taxon of <i>Galapagos</i> Tortoise. <i>PLoS ONE</i> , 2009, 4, e6272.	1.1	34
53	Patterns, Mechanisms and Genetics of Speciation in Reptiles and Amphibians. <i>Genes</i> , 2019, 10, 646.	1.0	33
54	Using digital images to reconstruct three-dimensional biological forms: a new tool for morphological studies. <i>Biological Journal of the Linnean Society</i> , 0, 95, 425-436.	0.7	32

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55	Genetic diversity and population structure of <i>Glossina pallidipes</i> in Uganda and western Kenya. <i>Parasites and Vectors</i> , 2011, 4, 122.	1.0	32
56	Multiple Origins of Cytologically Identical Chromosome Inversions in the <i>Anopheles gambiae</i> Complex. <i>Genetics</i> , 1998, 150, 807-814.	1.2	31
57	Population genomics through time provides insights into the consequences of decline and rapid demographic recovery through head-starting in a Galapagos giant tortoise. <i>Evolutionary Applications</i> , 2018, 11, 1811-1821.	1.5	29
58	The origin of captive Galapagos tortoises based on DNA analysis: implications for the management of natural populations. <i>Animal Conservation</i> , 2003, 6, 329-337.	1.5	28
59	Naturally rare versus newly rare: demographic inferences on two timescales inform conservation of Galapagos giant tortoises. <i>Ecology and Evolution</i> , 2015, 5, 676-694.	0.8	28
60	Theory, practice, and conservation in the age of genomics: The Galapagos giant tortoise as a case study. <i>Evolutionary Applications</i> , 2018, 11, 1084-1093.	1.5	28
61	Vectors as Epidemiological Sentinels: Patterns of Within-Tick <i>Borrelia burgdorferi</i> Diversity. <i>PLoS Pathogens</i> , 2016, 12, e1005759.	2.1	28
62	Temporal stability of <i>Glossina fuscipes fuscipes</i> populations in Uganda. <i>Parasites and Vectors</i> , 2011, 4, 19.	1.0	27
63	The population structure of <i>Glossina fuscipes fuscipes</i> in the Lake Victoria basin in Uganda: implications for vector control. <i>Parasites and Vectors</i> , 2012, 5, 222.	1.0	27
64	Identification of Genetically Important Individuals of the Rediscovered Floreana Galapagos Giant Tortoise ( <i>Chelonoidis elephantopus</i> ) Provides Founders for Species Restoration Program. <i>Scientific Reports</i> , 2017, 7, 11471.	1.6	27
65	Self-righting potential and the evolution of shell shape in Galapagos tortoises. <i>Scientific Reports</i> , 2017, 7, 15828.	1.6	27
66	Multiple evolutionary origins of <i>Trypanosoma evansi</i> in Kenya. <i>PLoS Neglected Tropical Diseases</i> , 2017, 11, e0005895.	1.3	27
67	A machine-learning approach to map landscape connectivity in <i>Aedes aegypti</i> with genetic and environmental data. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	3.3	27
68	Genetic diversity and population structure of the tsetse fly <i>Glossina fuscipes fuscipes</i> (Diptera: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 22). <i>Parasites and Vectors</i> , 2017, 11, e0005485.	1.3	26
69	<i>Wolbachia</i> association with the tsetse fly, <i>Glossina fuscipes fuscipes</i> , reveals high levels of genetic diversity and complex evolutionary dynamics. <i>BMC Evolutionary Biology</i> , 2013, 13, 31.	3.2	25
70	Genetic Diversity and Population Structure of <i>Trypanosoma brucei</i> in Uganda: Implications for the Epidemiology of Sleeping Sickness and Nagana. <i>PLoS Neglected Tropical Diseases</i> , 2015, 9, e0003353.	1.3	25
71	Complex interplay of evolutionary forces shaping population genomic structure of invasive <i>Aedes albopictus</i> in southern Europe. <i>PLoS Neglected Tropical Diseases</i> , 2019, 13, e0007554.	1.3	25
72	Giant Galapagos tortoises; molecular genetic analyses identify a trans-island hybrid in a repatriation program of an endangered taxon. <i>BMC Ecology</i> , 2007, 7, 2.	3.0	22

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73	Comparative genomics of drug resistance in <i>Trypanosoma brucei rhodesiense</i> . <i>Cellular and Molecular Life Sciences</i> , 2016, 73, 3387-3400.	2.4	22
74	Genome-Wide Assessment of Diversity and Divergence Among Extant Galapagos Giant Tortoise Species. <i>Journal of Heredity</i> , 2018, 109, 611-619.	1.0	22
75	Spatio-temporal distribution of <i>Spiroplasma</i> infections in the tsetse fly ( <i>Glossina fuscipes fuscipes</i> ) in northern Uganda. <i>PLoS Neglected Tropical Diseases</i> , 2019, 13, e0007340.	1.3	22
76	Development of new microsatellite loci and evaluation of loci from other pinniped species for the Galapagos sea lion ( <i>Zalophus californianus wollebaeki</i> ). <i>Conservation Genetics</i> , 2006, 7, 461-465.	0.8	21
77	De Novo Genome Assembly Shows Genome Wide Similarity between <i>Trypanosoma brucei brucei</i> and <i>Trypanosoma brucei rhodesiense</i> . <i>PLoS ONE</i> , 2016, 11, e0147660.	1.1	21
78	Cryptic east-west divergence and molecular diagnostics for two species of silver flies (Diptera: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 54) woolly adelgid. <i>Biological Control</i> , 2018, 121, 23-29.	1.4	20
79	Characterization of di-, tri- and tetranucleotide microsatellite markers with perfect repeats for <i>Trypanosoma brucei</i> and related species. <i>Molecular Ecology Notes</i> , 2006, 6, 508-510.	1.7	19
80	Genetics of a head-start program to guide conservation of an endangered Galapagos tortoise ( <i>Chelonoidis ephippium</i> ). <i>Conservation Genetics</i> , 2015, 16, 823-832.	0.8	18
81	Genetically informed captive breeding of hybrids of an extinct species of Galapagos tortoise. <i>Conservation Biology</i> , 2019, 33, 1404-1414.	2.4	18
82	Genetically Distinct <i>Glossina fuscipes fuscipes</i> Populations in the Lake Kyoga Region of Uganda and Its Relevance for Human African Trypanosomiasis. <i>BioMed Research International</i> , 2013, 2013, 1-12.	0.9	17
83	I-HEDGE: determining the optimum complementary sets of taxa for conservation using evolutionary isolation. <i>PeerJ</i> , 2016, 4, e2350.	0.9	17
84	Lineage Identification and Genealogical Relationships Among Captive Galapagos Tortoises. <i>Zoo Biology</i> , 2012, 31, 107-120.	0.5	16
85	<i>Babesia microti</i> from humans and ticks hold a genomic signature of strong population structure in the United States. <i>BMC Genomics</i> , 2016, 17, 888.	1.2	15
86	Temporal genetic differentiation in <i>Glossina pallidipes</i> tsetse fly populations in Kenya. <i>Parasites and Vectors</i> , 2017, 10, 471.	1.0	14
87	Colonization history of Galapagos giant tortoises: Insights from mitogenomes support the progression rule. <i>Journal of Zoological Systematics and Evolutionary Research</i> , 2020, 58, 1262-1275.	0.6	14
88	Evidence of temporal stability in allelic and mitochondrial haplotype diversity in populations of <i>Glossina fuscipes fuscipes</i> (Diptera: Glossinidae) in northern Uganda. <i>Parasites and Vectors</i> , 2016, 9, 258.	1.0	13
89	Multiple Paternity in the Norway Rat, <i>Rattus norvegicus</i> , from Urban Slums in Salvador, Brazil. <i>Journal of Heredity</i> , 2016, 107, 181-186.	1.0	13
90	Genomic analyses of African Trypanozoon strains to assess evolutionary relationships and identify markers for strain identification. <i>PLoS Neglected Tropical Diseases</i> , 2017, 11, e0005949.	1.3	13

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91	Patterns of Genome-Wide Variation in <i>Glossina fuscipes fuscipes</i> Tsetse Flies from Uganda. <i>G3: Genes, Genomes, Genetics</i> , 2016, 6, 1573-1584.	0.8	12
92	Whole genome sequencing shows sleeping sickness relapse is due to parasite regrowth and not reinfection. <i>Evolutionary Applications</i> , 2016, 9, 381-393.	1.5	12
93	Temporal Mitogenomics of the Galapagos Giant Tortoise from Pinzn Reveals Potential Biases in Population Genetic Inference. <i>Journal of Heredity</i> , 2018, 109, 631-640.	1.0	12
94	Genetic Pedigree Analysis of the Pilot Breeding Program for the Rediscovered Galapagos Giant Tortoise from Floreana Island. <i>Journal of Heredity</i> , 2018, 109, 620-630.	1.0	11
95	The population genomics of multiple tsetse fly ( <i>Glossina fuscipes fuscipes</i> ) admixture zones in Uganda. <i>Molecular Ecology</i> , 2019, 28, 66-85.	2.0	11
96	Restoration-mediated secondary contact leads to introgression of alewife ecotypes separated by a colonial-era dam. <i>Evolutionary Applications</i> , 2020, 13, 652-664.	1.5	10
97	Species delimitation and invasion history of the balsam woolly adelgid, <i>Adelges</i> ( <i>Dreyfusia</i> ) <i>piceae</i> (Hemiptera: Aphidoidea: Adelgidae), species complex. <i>Systematic Entomology</i> , 2021, 46, 186-204.	1.7	10
98	Habitat fragmentation and the genetic structure of the Amazonian palm <i>Mauritia flexuosa</i> L.f. (Arecaceae) on the island of Trinidad. <i>Conservation Genetics</i> , 2014, 15, 355-362.	0.8	9
99	Ecological and evolutionary influences on body size and shape in the Galpagos marine iguana ( <i>Amblyrhynchus cristatus</i> ). <i>Oecologia</i> , 2016, 181, 885-894.	0.9	9
100	Significant Genetic Impacts Accompany an Urban Rat Control Campaign in Salvador, Brazil. <i>Frontiers in Ecology and Evolution</i> , 2019, 7, .	1.1	9
101	Identification of winter moth ( <i>Operophtera brumata</i> ) refugia in North Africa and the Italian Peninsula during the last glacial maximum. <i>Ecology and Evolution</i> , 2019, 9, 13931-13941.	0.8	9
102	Uncovering Genomic Regions Associated with <i>Trypanosoma</i> Infections in Wild Populations of the Tsetse Fly <i>Glossina fuscipes</i> . <i>G3: Genes, Genomes, Genetics</i> , 2018, 8, 887-897.	0.8	8
103	A spatial genetics approach to inform vector control of tsetse flies ( <i>Glossina fuscipes</i> ) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50	0.8	8
104	Widespread hybridization among native and invasive species of Operophtera moths (Lepidoptera:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50	1.2	8
105	Genetic Differentiation of <i>Glossina pallidipes</i> Tsetse Flies in Southern Kenya. <i>American Journal of Tropical Medicine and Hygiene</i> , 2018, 99, 945-953.	0.6	8
106	Mitochondrial DNA sequence divergence and diversity of <i>Glossina fuscipes fuscipes</i> in the Lake Victoria basin of Uganda: implications for control. <i>Parasites and Vectors</i> , 2015, 8, 385.	1.0	7
107	Postglacial recolonization shaped the genetic diversity of the winter moth ( <i>Operophtera brumata</i> ) in Europe. <i>Ecology and Evolution</i> , 2017, 7, 3312-3323.	0.8	7
108	Demographic history and patterns of molecular evolution from whole genome sequencing in the radiation of Galapagos giant tortoises. <i>Molecular Ecology</i> , 2021, 30, 6325-6339.	2.0	7

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109	A set of highly discriminating microsatellite loci for the Galapagos marine iguana <i>Amblyrhynchus cristatus</i> . <i>Molecular Ecology Notes</i> , 2006, 6, 927-929.	1.7	6
110	Isolation of 13 novel highly polymorphic microsatellite loci for the Amazonian Palm <i>Mauritia flexuosa</i> L.f. (Arecaceae). <i>Conservation Genetics Resources</i> , 2012, 4, 355-357.	0.4	6
111	Potential arms race in the coevolution of primates and angiosperms: brazzein sweet proteins and gorilla taste receptors. <i>American Journal of Physical Anthropology</i> , 2016, 161, 181-185.	2.1	6
112	Phylogeography and population structure of the tsetse fly <i>Glossina pallidipes</i> in Kenya and the Serengeti ecosystem. <i>PLoS Neglected Tropical Diseases</i> , 2020, 14, e0007855.	1.3	6
113	A machine learning approach to integrating genetic and ecological data in tsetse flies ( <i>Glossina</i> ). <i>Trends in Ecology and Evolution</i> , 2021, 32, 1762-1777.	1.5	6
114	Genetic diversity of <i>Glossina fuscipes fuscipes</i> along the shores of Lake Victoria in Tanzania and Kenya: implications for management. <i>Parasites and Vectors</i> , 2017, 10, 268.	1.0	5
115	Was Frozen Mammoth or Giant Ground Sloth Served for Dinner at The Explorers Club?. <i>PLoS ONE</i> , 2016, 11, e0146825.	1.1	4
116	Seeking compromise across competing goals in conservation translocations: The case of the "extinct" Floreana Island Galapagos giant tortoise. <i>Journal of Applied Ecology</i> , 2020, 57, 136-148.	1.9	3
117	Evolution and phylogenetics. , 2021, , 117-138.		3
118	Northern Fennoscandia via the British Isles: evidence for a novel post-glacial recolonization route by winter moth ( <i>Operophtera brumata</i> ). <i>Frontiers of Biogeography</i> , 2021, 13, .	0.8	3
119	Four times out of Europe: Serial invasions of the winter moth, <i>Operophtera brumata</i> , to North America. <i>Molecular Ecology</i> , 2021, 30, 3439-3452.	2.0	3
120	A new lineage of Galapagos giant tortoises identified from museum samples. <i>Heredity</i> , 2022, 128, 261-270.	1.2	3
121	The Galapagos giant tortoise <i>Chelonoidis phantasticus</i> is not extinct. <i>Communications Biology</i> , 2022, 5, .	2.0	3
122	Real-time geographic settling of a hybrid zone between the invasive winter moth ( <i>Operophtera</i> ). <i>Evolutionary Ecology</i> , 2022, 36, 6617-6633.	2.0	2
123	Characterization of polymorphic microsatellite loci for the polychaete tubeworm <i>Hobsonia florida</i> . <i>Molecular Ecology Notes</i> , 2006, 6, 390-392.	1.7	1