

Andrew G Hope

List of Publications by Year in descending order

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Version: 2024-02-01

37
papers

1,292
citations

304743

22
h-index

361022

35
g-index

39
all docs

39
docs citations

39
times ranked

1370
citing authors

#	ARTICLE	IF	CITATIONS
1	Population genomics of free-ranging Great Plains white-tailed and mule deer reflects a long history of interspecific hybridization. <i>Evolutionary Applications</i> , 2022, 15, 111-131.	3.1	10
2	Misinterpretation of Genomic Data Matters for Endangered Species Listing: The Sub-specific Status of the Peñasco Least Chipmunk (<i>Neotamias minimus atristriatus</i>). <i>Frontiers in Conservation Science</i> , 2022, 2, .	1.9	2
3	Which mammals can be identified from camera traps and crowdsourced photographs?. <i>Journal of Mammalogy</i> , 2022, 103, 767-775.	1.3	12
4	Origins and diversity of the Bering Sea Island fauna: shifting linkages across the northern continents. <i>Biodiversity and Conservation</i> , 2021, 30, 1205-1232.	2.6	5
5	Consumer roles of small mammals within fragmented native tallgrass prairie. <i>Ecosphere</i> , 2021, 12, e03441.	2.2	7
6	Arctic Tundra Mammals. , 2020, , 356-373.		3
7	Harmony on the prairie? Grassland plant and animal community responses to variation in climate across land-use gradients. <i>Ecology</i> , 2020, 101, e02986.	3.2	16
8	Building an integrated infrastructure for exploring biodiversity: field collections and archives of mammals and parasites. <i>Journal of Mammalogy</i> , 2019, 100, 382-393.	1.3	61
9	Method for the Rapid Fixation of Gastrointestinal Helminths in Small Mammals. <i>Acta Parasitologica</i> , 2019, 64, 406-410.	1.1	4
10	Collection of Scientific Specimens: Benefits for Biodiversity Sciences and Limited Impacts on Communities of Small Mammals. <i>BioScience</i> , 2018, 68, 35-42.	4.9	32
11	Museum metabarcoding: A novel method revealing gut helminth communities of small mammals across space and time. <i>International Journal for Parasitology</i> , 2018, 48, 1061-1070.	3.1	26
12	The Beringian Coevolution Project: holistic collections of mammals and associated parasites reveal novel perspectives on evolutionary and environmental change in the North. <i>Arctic Science</i> , 2017, 3, 585-617.	2.3	50
13	Are the western water shrew (<i>Sorex navigator</i>) and American water shrew (<i>Sorex palustris</i>) morphologically distinct?. <i>Canadian Journal of Zoology</i> , 2017, 95, 727-736.	1.0	2
14	The Role of Temperature in the Distribution of the Glacier Ice Worm, <i>Mesenchytraeus solifugus</i> (Annelida: Oligochaeta: Enchytraeidae). <i>Arctic, Antarctic, and Alpine Research</i> , 2016, 48, 199-211.	1.1	20
15	Revision of widespread red squirrels (genus: <i>Tamiasciurus</i>) highlights the complexity of speciation within North American forests. <i>Molecular Phylogenetics and Evolution</i> , 2016, 100, 170-182.	2.7	59
16	Implications of the Circumpolar Genetic Structure of Polar Bears for Their Conservation in a Rapidly Warming Arctic. <i>PLoS ONE</i> , 2015, 10, e112021.	2.5	46
17	Arctic biodiversity: increasing richness accompanies shrinking refugia for a cold-associated tundra fauna. <i>Ecosphere</i> , 2015, 6, 1-67.	2.2	34
18	A multilocus evaluation of ermine (<i>Mustela erminea</i>) across the Holarctic, testing hypotheses of Pleistocene diversification in response to climate change. <i>Journal of Biogeography</i> , 2014, 41, 464-475.	3.0	32

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19	Multilocus phylogeography and systematic revision of North American water shrews (genus: <i>Sorex</i>). <i>Journal of Mammalogy</i> , 2014, 95, 722-738.	1.3	16
20	ACCOUNTING FOR RATE VARIATION AMONG LINEAGES IN COMPARATIVE DEMOGRAPHIC ANALYSES. <i>Evolution; International Journal of Organic Evolution</i> , 2014, 68, 2689-2700.	2.3	25
21	Development and characterization of 21 polymorphic microsatellite markers for the barren-ground shrew, <i>Sorex ugyunak</i> (Mammalia: Soricidae), through next-generation sequencing, and cross-species amplification in the masked shrew, <i>S. cinereus</i> . <i>Conservation Genetics Resources</i> , 2013, 5, 315-318.	0.8	4
22	Future distribution of tundra refugia in northern Alaska. <i>Nature Climate Change</i> , 2013, 3, 931-938.	18.8	34
23	Powassan Virus in Mammals, Alaska and New Mexico, USA, and Russia, 2004–2007. <i>Emerging Infectious Diseases</i> , 2013, 19, 2012-2016.	4.3	52
24	Temporal, spatial and ecological dynamics of speciation among amphibian small mammals. <i>Journal of Biogeography</i> , 2013, 40, 415-429.	3.0	34
25	High Shrew Diversity on Alaska's Seward Peninsula: Community Assembly and Environmental Change. <i>Northwestern Naturalist</i> , 2012, 93, 101-110.	0.4	6
26	A climate for speciation: Rapid spatial diversification within the <i>Sorex cinereus</i> complex of shrews. <i>Molecular Phylogenetics and Evolution</i> , 2012, 64, 671-684.	2.7	41
27	Shared Ancestry between a Newfound Mole-Borne Hantavirus and Hantaviruses Harbored by Cricetid Rodents. <i>Journal of Virology</i> , 2011, 85, 7496-7503.	3.4	71
28	Persistence and diversification of the Holarctic shrew, <i>Sorex tundrensis</i> (Family Soricidae), in response to climate change. <i>Molecular Ecology</i> , 2011, 20, 4346-4370.	3.9	30
29	Novel Hantavirus in the Flat-Skulled Shrew (<i>Sorex roboratus</i>). <i>Vector-Borne and Zoonotic Diseases</i> , 2010, 10, 593-597.	1.5	44
30	High-latitude diversification within Eurasian least shrews and Alaska tiny shrews (Soricidae). <i>Journal of Mammalogy</i> , 2010, 91, 1041-1057.	1.3	44
31	Evolutionary Insights from a Genetically Divergent Hantavirus Harbored by the European Common Mole (<i>Talpa europaea</i>). <i>PLoS ONE</i> , 2009, 4, e6149.	2.5	107
32	Genetic diversity and phylogeography of Seewis virus in the Eurasian common shrew in Finland and Hungary. <i>Virology Journal</i> , 2009, 6, 208.	3.4	45
33	Phylogenetically Distinct Hantaviruses in the Masked Shrew (<i>Sorex cinereus</i>) and Dusky Shrew (<i>Sorex</i>) Tj ETQq1 1 Q.784314 ggBT /Over	1.4	84
34	Phylogenetically distinct hantaviruses in the masked shrew (<i>Sorex cinereus</i>) and dusky shrew (<i>Sorex</i>) Tj ETQq0 0 0 ggBT /Over lock 10 Tf	1.4	45
35	Beringia: Intercontinental exchange and diversification of high latitude mammals and their parasites during the Pliocene and Quaternary. <i>Mammal Study</i> , 2005, 30, S33-S44.	0.6	81
36	Satellite imagery characterizes local animal reservoir populations of Sin Nombre virus in the southwestern United States. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002, 99, 16817-16822.	7.1	103

#	ARTICLE	IF	CITATIONS
37	Speciation of North American pygmy shrews (Eulipotyphla: Soricidae) supports spatial but not temporal congruence of diversification among boreal species. <i>Biological Journal of the Linnean Society</i> , 0, , .	1.6	2