Michael V Westbury

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Evaluating the role of referenceâ€genome phylogenetic distance on evolutionary inference. Molecular Ecology Resources, 2022, 22, 45-55.	2.2	28
2	Ancient Mitogenomes Suggest Stable Mitochondrial Clades of the Siberian Roe Deer. Genes, 2022, 13, 114.	1.0	3
3	High genomic diversity in the endangered East Greenland Svalbard Barents Sea stock of bowhead whales (Balaena mysticetus). Scientific Reports, 2022, 12, 6118.	1.6	2
4	Grey wolf genomic history reveals a dual ancestry of dogs. Nature, 2022, 607, 313-320.	13.7	48
5	Identifying the true number of specimens of the extinct blue antelope (Hippotragus leucophaeus). Scientific Reports, 2021, 11, 2100.	1.6	9
6	Ancient mitochondrial genomes from Chinese cave hyenas provide insights into the evolutionary history of the genus <i>Crocuta</i> . Proceedings of the Royal Society B: Biological Sciences, 2021, 288, 20202934.	1.2	9
7	Ecological Specialization and Evolutionary Reticulation in Extant Hyaenidae. Molecular Biology and Evolution, 2021, 38, 3884-3897.	3.5	15
8	Ocean-wide genomic variation in Gray's beaked whales, <i>Mesoplodon grayi</i> . Royal Society Open Science, 2021, 8, 201788.	1.1	11
9	Genomic consequences of humanâ€mediated translocations in margin populations of an endangered amphibian. Evolutionary Applications, 2021, 14, 1623-1634.	1.5	2
10	Circumpolar phylogeography and demographic history of beluga whales reflect past climatic fluctuations. Molecular Ecology, 2021, 30, 2543-2559.	2.0	12
11	African and Asian leopards are highly differentiated at the genomic level. Current Biology, 2021, 31, 1872-1882.e5.	1.8	20
12	Southern introgression increases adaptive immune gene variability in northern range margin populations of Fireâ€bellied toad. Ecology and Evolution, 2021, 11, 9776-9790.	0.8	2
13	A genomic exploration of the early evolution of extant cats and their sabre-toothed relatives. Open Research Europe, 2021, 1, 25.	2.0	2
14	Successful application of ancient DNA extraction and library construction protocols to museum wet collection specimens. Molecular Ecology Resources, 2021, 21, 2299-2315.	2.2	36
15	Historical population declines prompted significant genomic erosion in the northern and southern white rhinoceros (<i>Ceratotherium simum</i>). Molecular Ecology, 2021, 30, 6355-6369.	2.0	39
16	A sliver of the past: The decimation of the genetic diversity of the Mexican wolf. Molecular Ecology, 2021, 30, 6340-6354.	2.0	6
17	Diversity and Paleodemography of the Addax (Addax nasomaculatus), a Saharan Antelope on the Verge of Extinction. Genes, 2021, 12, 1236.	1.0	8
18	Ancient and modern genomes unravel the evolutionary history of the rhinoceros family. Cell, 2021, 184, 4874-4885.e16.	13.5	49

MICHAEL V WESTBURY

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19	Genomic Adaptations and Evolutionary History of the Extinct Scimitar-Toothed Cat, Homotherium latidens. Current Biology, 2020, 30, 5018-5025.e5.	1.8	34
20	Analyses of key genes involved in Arctic adaptation in polar bears suggest selection on both standing variation and de novo mutations played an important role. BMC Genomics, 2020, 21, 543.	1.2	3
21	Palaeoproteomic analysis of Pleistocene cave hyenas from east Asia. Scientific Reports, 2020, 10, 16674.	1.6	4
22	Complete mitochondrial genomes offer insights into the evolutionary relationships and comparative genetic diversity of New Zealand's iconic kiwi (<i>Apteryx</i> spp.). New Zealand Journal of Zoology, 2020, 47, 291-299.	0.6	2
23	Hyena paleogenomes reveal a complex evolutionary history of cross-continental gene flow between spotted and cave hyena. Science Advances, 2020, 6, eaay0456.	4.7	38
24	Interspecific Gene Flow and the Evolution of Specialization in Black and White Rhinoceros. Molecular Biology and Evolution, 2020, 37, 3105-3117.	3.5	20
25	Genomic analyses reveal an absence of contemporary introgressive admixture between fin whales and blue whales, despite known hybrids. PLoS ONE, 2019, 14, e0222004.	1.1	15
26	Hybridization between two high Arctic cetaceans confirmed by genomic analysis. Scientific Reports, 2019, 9, 7729.	1.6	33
27	Paleogenome Reveals Genetic Contribution of Extinct Giant Panda to Extant Populations. Current Biology, 2019, 29, 1695-1700.e6.	1.8	22
28	The complete mitochondrial genome of a European fire-bellied toad (<i>Bombina bombina</i>) from Germany. Mitochondrial DNA Part B: Resources, 2019, 4, 498-500.	0.2	3
29	Narwhal Genome Reveals Long-Term Low Genetic Diversity despite Current Large Abundance Size. IScience, 2019, 15, 592-599.	1.9	49
30	Aardwolf Population Diversity and Phylogenetic Positioning Inferred Using Complete Mitochondrial Genomes. African Journal of Wildlife Research, 2019, 49, .	0.2	0
31	Cryptic species in a well-known habitat: applying taxonomics to the amphipod genus Epimeria (Crustacea, Peracarida). Scientific Reports, 2018, 8, 6893.	1.6	15
32	The complete mitochondrial genome of the common vole, Microtus arvalis (Rodentia: Arvicolinae). Mitochondrial DNA Part B: Resources, 2018, 3, 446-447.	0.2	9
33	Extended and Continuous Decline in Effective Population Size Results in Low Genomic Diversity in the World's Rarest Hyena Species, the Brown Hyena. Molecular Biology and Evolution, 2018, 35, 1225-1237.	3.5	72
34	Ancient DNA from Giant Panda (Ailuropoda melanoleuca) of South-Western China Reveals Genetic Diversity Loss during the Holocene. Genes, 2018, 9, 198.	1.0	14
35	Evolutionary History of Saber-Toothed Cats Based on Ancient Mitogenomics. Current Biology, 2017, 27, 3330-3336.e5.	1.8	45
36	Reduction of the contaminant fraction of DNA obtained from an ancient giant panda bone. BMC Research Notes, 2017, 10, 754.	0.6	26

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37	A genomic exploration of the early evolution of extant cats and their sabre-toothed relatives. Open Research Europe, 0, 1, 25.	2.0	1
38	Palaeogenome Reveals Genetic Contribution of Extinct Giant Panda to Extant Populations. SSRN Electronic Journal, 0, , .	0.4	0