Erwan Quéméré

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Landscape genetics of an endangered lemur (<i>Propithecus tattersalli</i>) within its entire fragmented range. Molecular Ecology, 2010, 19, 1606-1621.	3.9	156
2	A DNA Metabarcoding Study of a Primate Dietary Diversity and Plasticity across Its Entire Fragmented Range. PLoS ONE, 2013, 8, e58971.	2.5	89
3	Genetic data suggest a natural prehuman origin of open habitats in northern Madagascar and question the deforestation narrative in this region. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 13028-13033.	7.1	81
4	Molecular phylogeny in mytilids supports the wooden steps to deep-sea vents hypothesis. Comptes Rendus - Biologies, 2007, 330, 446-456.	0.2	64
5	Climate change and human colonization triggered habitat loss and fragmentation in Madagascar. Molecular Ecology, 2017, 26, 5203-5222.	3.9	56
6	Immunogenetic heterogeneity in a widespread ungulate: the European roe deer (<i>Capreolu</i> s) Tj ETQq0 0 0	rg <u>B</u> Ţ/Ove	rlo <u>ç</u> k 10 Tf 5(
7	An improved PCR-based method for faster sex determination in brown trout (Salmo trutta) and Atlantic salmon (Salmo salar). Conservation Genetics Resources, 2014, 6, 825-827.	0.8	34
8	Non-invasive conservation genetics of the critically endangered golden-crowned sifaka (Propithecus) Tj ETQq0 0 Genetics, 2010, 11, 675-687.	0 rgBT /Ov 1.5	verlock 10 Tf 28
9	Spatial variation in density and total size estimates in fragmented primate populations: the goldenâ€crowned sifaka (<i>Propithecus tattersalli</i>). American Journal of Primatology, 2010, 72, 72-80.	1.7	27
10	Seascape and its effect on migratory lifeâ€history strategy influences gene flow among coastal brown trout (<i>Salmo trutta</i>) populations in the English Channel. Journal of Biogeography, 2016, 43, 498-509.	3.0	18
11	RADâ€sequencing for estimating genomic relatedness matrixâ€based heritability in the wild: A case study in roe deer. Molecular Ecology Resources, 2019, 19, 1205-1217.	4.8	18
12	Genetic consequences of social structure in the golden-crowned sifaka. Heredity, 2020, 125, 328-339.	2.6	14
13	Pedigreeâ€free quantitative genetic approach provides evidence for heritability of movement tactics in wild roe deer. Journal of Evolutionary Biology, 2020, 33, 595-607.	1.7	14
14	Demography and adaptation promoting evolutionary transitions in a mammalian genus that diversified during the Pleistocene. Molecular Ecology, 2020, 29, 2777-2792.	3.9	13
15	Pathogenâ€mediated selection favours the maintenance of innate immunity gene polymorphism in a widespread wild ungulate. Journal of Evolutionary Biology, 2021, 34, 1156-1166.	1.7	13
16	Genetic epidemiology of the Alpine ibex reservoir of persistent and virulent brucellosis outbreak. Scientific Reports, 2020, 10, 4400.	3.3	12
17	Between-population differences in the genetic and maternal components of body mass in roe deer. BMC Evolutionary Biology, 2018, 18, 39.	3.2	10
18	Unraveling the dietary diversity of Neotropical top predators using scat DNA metabarcoding: A case study on the elusive Giant Otter. Environmental DNA, 2021, 3, 889-900.	5.8	8

#	Article	IF	CITATIONS
19	Reduced microsatellite heterozygosity does not affect natal dispersal in three contrasting roe deer populations. Oecologia, 2015, 177, 631-643.	2.0	5
20	Immune gene variability influences roe deer natal dispersal. Oikos, 2016, 125, 1790-1801.	2.7	5
21	Urine DNA (uDNA) as a nonâ€lethal method for endoparasite biomonitoring: Development and validation. Environmental DNA, 2021, 3, 1035-1045.	5.8	5
22	Quantifying heritability and estimating evolutionary potential in the wild when individuals that share genes also share environments. Journal of Animal Ecology, 2022, 91, 1239-1250.	2.8	5
23	Characterization of 20 microsatellites marker loci in the golden-crowned sifaka (Propithecus) Tj ETQq1 1 0.7843	14 _{.1} gBT /C	Verlock 10
24	Maternal effects shape offspring physiological condition but do not senesce in a wild mammal. Journal of Evolutionary Biology, 2021, 34, 661-670.	1.7	1