

Laurent Frantz

List of Publications by Year in descending order

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Version: 2024-02-01

17
papers

845
citations

687220

13
h-index

887953

17
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18
all docs

18
docs citations

18
times ranked

1436
citing authors

#	ARTICLE	IF	CITATIONS
1	Natural and human-driven selection of a single non-coding body size variant in ancient and modern canids. <i>Current Biology</i> , 2022, 32, 889-897.e9.	1.8	23
2	Grey wolf genomic history reveals a dual ancestry of dogs. <i>Nature</i> , 2022, 607, 313-320.	13.7	48
3	Reconsidering domestication from a process archaeology perspective. <i>World Archaeology</i> , 2021, 53, 56-77.	0.5	36
4	Modern Siberian dog ancestry was shaped by several thousand years of Eurasian-wide trade and human dispersal. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	3.3	19
5	Ancient DNA suggests modern wolves trace their origin to a Late Pleistocene expansion from Beringia. <i>Molecular Ecology</i> , 2020, 29, 1596-1610.	2.0	70
6	Origins and genetic legacy of prehistoric dogs. <i>Science</i> , 2020, 370, 557-564.	6.0	152
7	Genomic consequences of population decline in critically endangered pangolins and their demographic histories. <i>National Science Review</i> , 2020, 7, 798-814.	4.6	45
8	Genetic Diversity, Inbreeding Level, and Genetic Load in Endangered Snub-Nosed Monkeys (<i>Rhinopithecus</i>). <i>Frontiers in Genetics</i> , 2020, 11, 615926.	1.1	14
9	Specialized sledge dogs accompanied Inuit dispersal across the North American Arctic. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2019, 286, 20191929.	1.2	38
10	The Origin and Population History of the Endangered Golden Snub-Nosed Monkey (<i>Rhinopithecus</i>)	3.5	20
11	Analysis of the genetic variation in mitochondrial DNA, Y-chromosome sequences, and MC1R sheds light on the ancestry of Nigerian indigenous pigs. <i>Genetics Selection Evolution</i> , 2017, 49, 52.	1.2	8
12	Reconstructing Asian faunal introductions to eastern Africa from multi-proxy biomolecular and archaeological datasets. <i>PLoS ONE</i> , 2017, 12, e0182565.	1.1	53
13	A novel MC1R allele for black coat colour reveals the Polynesian ancestry and hybridization patterns of Hawaiian feral pigs. <i>Royal Society Open Science</i> , 2016, 3, 160304.	1.1	19
14	The Evolution of Suidae. <i>Annual Review of Animal Biosciences</i> , 2016, 4, 61-85.	3.6	85
15	The genetics of indirect ecological effects: plant parasites and aphid herbivores. <i>Frontiers in Genetics</i> , 2014, 5, 72.	1.1	2
16	Host-plant genotypic diversity and community genetic interactions mediate aphid spatial distribution. <i>Ecology and Evolution</i> , 2014, 4, 121-131.	0.8	12
17	Pig Domestication and Human-Mediated Dispersal in Western Eurasia Revealed through Ancient DNA and Geometric Morphometrics. <i>Molecular Biology and Evolution</i> , 2013, 30, 824-832.	3.5	196