

Eleftheria Palkopoulou

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/11386949/publications.pdf>

Version: 2024-02-01

15
papers

1,187
citations

759233

12
h-index

996975

15
g-index

16
all docs

16
docs citations

16
times ranked

2194
citing authors

#	ARTICLE	IF	CITATIONS
1	Ancient Wolf Genome Reveals an Early Divergence of Domestic Dog Ancestors and Admixture into High-Latitude Breeds. <i>Current Biology</i> , 2015, 25, 1515-1519.	3.9	270
2	Complete Genomes Reveal Signatures of Demographic and Genetic Declines in the Woolly Mammoth. <i>Current Biology</i> , 2015, 25, 1395-1400.	3.9	263
3	A comprehensive genomic history of extinct and living elephants. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E2566-E2574.	7.1	142
4	Early Pleistocene enamel proteome from Dmanisi resolves <i>Stephanorhinus</i> phylogeny. <i>Nature</i> , 2019, 574, 103-107.	27.8	135
5	Serial population extinctions in a small mammal indicate Late Pleistocene ecosystem instability. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 20532-20536.	7.1	80
6	Holarctic genetic structure and range dynamics in the woolly mammoth. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2013, 280, 20131910.	2.6	72
7	Palaeogenomes of Eurasian straight-tusked elephants challenge the current view of elephant evolution. <i>ELife</i> , 2017, 6, .	6.0	50
8	Synchronous genetic turnovers across Western Eurasia in Late Pleistocene collared lemmings. <i>Global Change Biology</i> , 2016, 22, 1710-1721.	9.5	45
9	Resolution of the type material of the Asian elephant, <i>Elephas maximus</i> Linnaeus, 1758 (Proboscidea, Elephantidae). <i>Zoological Journal of the Linnean Society</i> , 2014, 170, 222-232.	2.3	31
10	Resolution of the type material of the Asian elephant, <i>Elephas maximus</i> Linnaeus, 1758 (Proboscidea,)	2.9	22
11	Mitogenome evolution in the last surviving woolly mammoth population reveals neutral and functional consequences of small population size. <i>Evolution Letters</i> , 2017, 1, 292-303.	3.3	22
12	Competitive mapping allows for the identification and exclusion of human DNA contamination in ancient faunal genomic datasets. <i>BMC Genomics</i> , 2020, 21, 844.	2.8	15
13	Population structure and recent temporal changes in genetic variation in Eurasian otters from Sweden. <i>Conservation Genetics</i> , 2015, 16, 371-384.	1.5	14
14	Genetic signs of multiple colonization events in Baltic ciscoes with radiation into sympatric spring and autumn spawners confined to early postglacial arrival. <i>Ecology and Evolution</i> , 2014, 4, 4346-4360.	1.9	13
15	Back to BaySICS: A User-Friendly Program for Bayesian Statistical Inference from Coalescent Simulations. <i>PLoS ONE</i> , 2014, 9, e98011.	2.5	11