Eleftheria Palkopoulou

List of Publications by Year
in descending order

15	1,187	759233 12	996975 15
papers	citations	h-index	g-index

16	16	16	2194
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Competitive mapping allows for the identification and exclusion of human DNA contamination in ancient faunal genomic datasets. BMC Genomics, 2020, 21, 844.	2.8	15
2	Early Pleistocene enamel proteome from Dmanisi resolves Stephanorhinus phylogeny. Nature, 2019, 574, 103-107.	27.8	135
3	A comprehensive genomic history of extinct and living elephants. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E2566-E2574.	7.1	142
4	Mitogenome evolution in the last surviving woolly mammoth population reveals neutral and functional consequences of small population size. Evolution Letters, 2017, 1, 292-303.	3.3	22
5	Palaeogenomes of Eurasian straight-tusked elephants challenge the current view of elephant evolution. ELife, 2017, 6, .	6.0	50
6	Synchronous genetic turnovers across Western Eurasia in Late Pleistocene collared lemmings. Global Change Biology, 2016, 22, 1710-1721.	9.5	45
7	Ancient Wolf Genome Reveals an Early Divergence of Domestic Dog Ancestors and Admixture into High-Latitude Breeds. Current Biology, 2015, 25, 1515-1519.	3.9	270
8	Complete Genomes Reveal Signatures of Demographic and Genetic Declines in the Woolly Mammoth. Current Biology, 2015, 25, 1395-1400.	3.9	263
9	Population structure and recent temporal changes in genetic variation in Eurasian otters from Sweden. Conservation Genetics, 2015, 16, 371-384.	1.5	14
10	Resolution of the type material of the Asian elephant,Elephas maximusLinnaeus, 1758 (Proboscidea, Elephantidae). Zoological Journal of the Linnean Society, 2014, 170, 222-232.	2.3	31
11	Resolution of the type material of the Asian elephant, Elephas maximus Linnaeus, 1758 (Proboscidea,) Tj ETQq1 1 0.784314 rgBT /Ov	2.3	22
12	Genetic signs of multiple colonization events in Baltic ciscoes with radiation into sympatric springâ€• and autumnâ€spawners confined to early postglacial arrival. Ecology and Evolution, 2014, 4, 4346-4360.	1.9	13
13	Back to BaySICS: A User-Friendly Program for Bayesian Statistical Inference from Coalescent Simulations. PLoS ONE, 2014, 9, e98011.	2.5	11
14	Holarctic genetic structure and range dynamics in the woolly mammoth. Proceedings of the Royal Society B: Biological Sciences, 2013, 280, 20131910.	2.6	72
15	Serial population extinctions in a small mammal indicate Late Pleistocene ecosystem instability. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 20532-20536.	7.1	80