

Gui-Lian Sheng

List of Publications by Year in descending order

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Version: 2024-02-01

18
papers

237
citations

1163117

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docs citations

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times ranked

456
citing authors

#	ARTICLE	IF	CITATIONS
1	Hyena paleogenomes reveal a complex evolutionary history of cross-continental gene flow between spotted and cave hyena. <i>Science Advances</i> , 2020, 6, eaay0456.	10.3	38
2	Once lost, twice found: Combined analysis of ancient giant panda sequences characterises extinct clade. <i>Journal of Biogeography</i> , 2019, 46, 251-253.	3.0	37
3	Pleistocene Chinese cave hyenas and the recent Eurasian history of the spotted hyena, <i>Crocuta crocuta</i> . <i>Molecular Ecology</i> , 2014, 23, 522-533.	3.9	29
4	Reduction of the contaminant fraction of DNA obtained from an ancient giant panda bone. <i>BMC Research Notes</i> , 2017, 10, 754.	1.4	26
5	Paleogenome Reveals Genetic Contribution of Extinct Giant Panda to Extant Populations. <i>Current Biology</i> , 2019, 29, 1695-1700.e6.	3.9	22
6	Molecular identification of late and terminal Pleistocene <i>Equus ovodovi</i> from northeastern China. <i>PLoS ONE</i> , 2019, 14, e0216883.	2.5	15
7	Ancient DNA from Giant Panda (<i>Ailuropoda melanoleuca</i>) of South-Western China Reveals Genetic Diversity Loss during the Holocene. <i>Genes</i> , 2018, 9, 198.	2.4	14
8	Deep genetic divergence within a living fossil brachiopod <i>Lingula anatina</i> . <i>Journal of Paleontology</i> , 2013, 87, 902-908.	0.8	10
9	Ancient DNA sequences from <i>Coelodonta antiquitatis</i> in China reveal its divergence and phylogeny. <i>Science China Earth Sciences</i> , 2014, 57, 388-396.	5.2	10
10	Mitochondrial genomes of Late Pleistocene caballine horses from China belong to a separate clade. <i>Quaternary Science Reviews</i> , 2020, 250, 106691.	3.0	9
11	Ancient mitochondrial genomes from Chinese cave hyenas provide insights into the evolutionary history of the genus <i>Crocuta</i> . <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2021, 288, 20202934.	2.6	9
12	DNA analyses of wild boar remains from archaeological sites in Guangxi, China. <i>Quaternary International</i> , 2014, 354, 147-153.	1.5	4
13	Different maternal lineages revealed by ancient mitochondrial genome of <i>Camelus bactrianus</i> from China. <i>Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis</i> , 2019, 30, 786-793.	0.7	4
14	Ancient DNA of northern China Hystricidae sub-fossils reveals the evolutionary history of old world porcupines in the Late Pleistocene. <i>BMC Evolutionary Biology</i> , 2020, 20, 88.	3.2	4
15	Ancient Mitogenomes Suggest Stable Mitochondrial Clades of the Siberian Roe Deer. <i>Genes</i> , 2022, 13, 114.	2.4	3
16	Ancient Mitogenomes Provide New Insights into the Origin and Early Introduction of Chinese Domestic Donkeys. <i>Frontiers in Genetics</i> , 2021, 12, 759831.	2.3	2
17	Short sequence effect of ancient DNA on mammoth phylogenetic analyses. <i>Frontiers of Earth Science</i> , 2009, 3, 100-106.	0.5	0
18	Palaeogenome Reveals Genetic Contribution of Extinct Giant Panda to Extant Populations. <i>SSRN Electronic Journal</i> , 0, , .	0.4	0