

Clio Der Sarkissian

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

34
papers

1,957
citations

21
h-index

34
g-index

34
ext. papers

2,694
ext. citations

8.6
avg, IF

4.06
L-index

#	Paper	IF	Citations
34	Sex in the city: Uncovering sex-specific management of equine resources from prehistoric times to the Modern Period in France. <i>Journal of Archaeological Science: Reports</i> , 2022 , 41, 103341	0.7	0
33	Historical management of equine resources in France from the Iron Age to the Modern Period. <i>Journal of Archaeological Science: Reports</i> , 2021 , 40, 103250	0.7	3
32	The origins and spread of domestic horses from the Western Eurasian steppes. <i>Nature</i> , 2021 , 598, 634-640.	40.4	24
31	Heterogeneous Hunter-Gatherer and Steppe-Related Ancestries in Late Neolithic and Bell Beaker Genomes from Present-Day France. <i>Current Biology</i> , 2021 , 31, 1072-1083.e10	6.3	5
30	No particular genomic features underpin the dramatic economic consequences of 17 century plague epidemics in Italy. <i>iScience</i> , 2021 , 24, 102383	6.1	4
29	Ancient horse genomes reveal the timing and extent of dispersals across the Bering Land Bridge. <i>Molecular Ecology</i> , 2021 , 30, 6144-6161	5.7	7
28	Assessing the predictive taxonomic power of the bony labyrinth 3D shape in horses, donkeys and their F1-hybrids. <i>Journal of Archaeological Science</i> , 2021 , 131, 105383	2.9	4
27	Ancient DNA analysis. <i>Nature Reviews Methods Primers</i> , 2021 , 1,		21
26	Ancient Metagenomic Studies: Considerations for the Wider Scientific Community.. <i>MSystems</i> , 2021 , 6, e0131521	7.6	
25	Unveiling the Ecological Applications of Ancient DNA From Mollusk Shells. <i>Frontiers in Ecology and Evolution</i> , 2020 , 8,	3.7	15
24	CASCADE: A Custom-Made Archiving System for the Conservation of Ancient DNA Experimental Data. <i>Frontiers in Ecology and Evolution</i> , 2020 , 8,	3.7	1
23	Displays an Original Type IV Secretion System in Strains Pathogenic for Bivalve Molluscs. <i>Frontiers in Microbiology</i> , 2018 , 9, 227	5.7	10
22	Ancient DNA analysis identifies marine mollusc shells as new metagenomic archives of the past. <i>Molecular Ecology Resources</i> , 2017 , 17, 835-853	8.4	38
21	Less effective selection leads to larger genomes. <i>Genome Research</i> , 2017 , 27, 1016-1028	9.7	44
20	Ancient genomic changes associated with domestication of the horse. <i>Science</i> , 2017 , 356, 442-445	33.3	149
19	Combining bleach and mild predigestion improves ancient DNA recovery from bones. <i>Molecular Ecology Resources</i> , 2017 , 17, 742-751	8.4	48
18	Evolutionary Patterns and Processes: Lessons from Ancient DNA. <i>Systematic Biology</i> , 2017 , 66, e1-e29	8.4	53

17	metaBIT, an integrative and automated metagenomic pipeline for analysing microbial profiles from high-throughput sequencing shotgun data. <i>Molecular Ecology Resources</i> , 2016 , 16, 1415-1427	8.4	30
16	The Evolutionary Origin and Genetic Makeup of Domestic Horses. <i>Genetics</i> , 2016 , 204, 423-434	4	39
15	Mitochondrial genomes reveal the extinct Hippidion as an outgroup to all living equids. <i>Biology Letters</i> , 2015 , 11,	3.6	26
14	Evolutionary Genomics and Conservation of the Endangered Przewalski's Horse. <i>Current Biology</i> , 2015 , 25, 2577-83	6.3	115
13	Tracking the origins of Yakutian horses and the genetic basis for their fast adaptation to subarctic environments. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015 , 112, E6889-97	11.5	89
12	Major transitions in human evolution revisited: a tribute to ancient DNA. <i>Journal of Human Evolution</i> , 2015 , 79, 4-20	3.1	35
11	Ancient and modern environmental DNA. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2015 , 370, 20130383	5.8	184
10	Pros and cons of methylation-based enrichment methods for ancient DNA. <i>Scientific Reports</i> , 2015 , 5, 11826	4.9	40
9	Ancient genomics. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2015 , 370, 20130387	3.8	107
8	Characterization of ancient and modern genomes by SNP detection and phylogenomic and metagenomic analysis using PALEOMIX. <i>Nature Protocols</i> , 2014 , 9, 1056-82	18.8	231
7	Rodents of the Caribbean: origin and diversification of hutias unravelled by next-generation museomics. <i>Biology Letters</i> , 2014 , 10,	3.6	71
6	Prehistoric genomes reveal the genetic foundation and cost of horse domestication. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014 , 111, E5661-9	11.5	197
5	Mitochondrial genome sequencing in Mesolithic North East Europe Unearths a new sub-clade within the broadly distributed human haplogroup C1. <i>PLoS ONE</i> , 2014 , 9, e87612	3.7	15
4	Ancient DNA reveals prehistoric gene-flow from siberia in the complex human population history of North East Europe. <i>PLoS Genetics</i> , 2013 , 9, e1003296	6	66
3	Neolithic mitochondrial haplogroup H genomes and the genetic origins of Europeans. <i>Nature Communications</i> , 2013 , 4, 1764	17.4	141
2	The influence of rate heterogeneity among sites on the time dependence of molecular rates. <i>Molecular Biology and Evolution</i> , 2012 , 29, 3345-58	8.3	140
1	Population history from the Neolithic to present on the Mediterranean island of Sardinia: An ancient DNA perspective		5