Ashot Margaryan

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3629229/publications.pdf

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30 4,982 21 papers citations h-index

39 39 39 6326 all docs docs citations times ranked citing authors

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g-index

#	Article	IF	CITATIONS
1	Population genomics of Bronze Age Eurasia. Nature, 2015, 522, 167-172.	27.8	1,166
2	A genomic history of Aboriginal Australia. Nature, 2016, 538, 207-214.	27.8	439
3	137 ancient human genomes from across the Eurasian steppes. Nature, 2018, 557, 369-374.	27.8	325
4	The prehistoric peopling of Southeast Asia. Science, 2018, 361, 88-92.	12.6	291
5	Genomic structure in Europeans dating back at least 36,200 years. Science, 2014, 346, 1113-1118.	12.6	287
6	Ancient genomes show social and reproductive behavior of early Upper Paleolithic foragers. Science, 2017, 358, 659-662.	12.6	263
7	The first horse herders and the impact of early Bronze Age steppe expansions into Asia. Science, 2018, 360, .	12.6	262
8	The population history of northeastern Siberia since the Pleistocene. Nature, 2019, 570, 182-188.	27.8	259
9	Dense sampling of bird diversity increases power of comparative genomics. Nature, 2020, 587, 252-257.	27.8	251
10	Early human dispersals within the Americas. Science, 2018, 362, .	12.6	230
11	Improving access to endogenous DNA in ancient bones and teeth. Scientific Reports, 2015, 5, 11184.	3.3	182
12	Population genomics of the Viking world. Nature, 2020, 585, 390-396.	27.8	143
13	Ancient genomics. Philosophical Transactions of the Royal Society B: Biological Sciences, 2015, 370, 20130387.	4.0	142
14	Comparing Ancient DNA Preservation in Petrous Bone and Tooth Cementum. PLoS ONE, 2017, 12, e0170940.	2.5	136
15	Unraveling ancestry, kinship, and violence in a Late Neolithic mass grave. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 10705-10710.	7.1	119
16	Diverse variola virus (smallpox) strains were widespread in northern Europe in the Viking Age. Science, 2020, 369, .	12.6	108
17	Ancient human parvovirus B19 in Eurasia reveals its long-term association with humans. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 7557-7562.	7.1	64
18	Ancient and modern genomes unravel the evolutionary history of the rhinoceros family. Cell, 2021, 184, 4874-4885.e16.	28.9	49

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19	Ancient pathogen <scp>DNA</scp> in human teeth and petrous bones. Ecology and Evolution, 2018, 8, 3534-3542.	1.9	38
20	Eight Millennia of Matrilineal Genetic Continuity in the South Caucasus. Current Biology, 2017, 27, 2023-2028.e7.	3.9	37
21	Screening archaeological bone for palaeogenetic and palaeoproteomic studies. PLoS ONE, 2020, 15, e0235146.	2.5	34
22	Evolutionary History, Genomic Adaptation to Toxic Diet, and Extinction of the Carolina Parakeet. Current Biology, 2020, 30, 108-114.e5.	3.9	24
23	Mitochondrial genomes of Danish vertebrate species generated for the national DNA reference database, DNAmark. Environmental DNA, 2021, 3, 472-480.	5.8	24
24	Recent mitochondrial lineage extinction in the critically endangered Javan rhinoceros. Zoological Journal of the Linnean Society, 2020, 190, 372-383.	2.3	13
25	Genomic Steppe ancestry in skeletons from the Neolithic Single Grave Culture in Denmark. PLoS ONE, 2021, 16, e0244872.	2.5	11
26	Regionalized autosomal STR profiles among Armenian groups suggest disparate genetic influences. American Journal of Physical Anthropology, 2011, 146, 171-178.	2.1	10
27	High Yâ€chromosomal Differentiation Among Ethnic Groups of Dir and Swat Districts, Pakistan. Annals of Human Genetics, 2017, 81, 234-248.	0.8	9
28	Uncovering the genomic and metagenomic research potential in old ethanol-preserved snakes. PLoS ONE, 2021, 16, e0256353.	2.5	6
29	Paternal Lineage Analysis Supports an Armenian Rather Than a Central Asian Genetic Origin of the Hamshenis. Human Biology, 2012, 84, 405-422.	0.2	1
30	The genomic origin of Zana of Abkhazia. Genetics & Genomics Next, 2021, 2, e10051.	1.5	0