

Monika Karmin

List of Publications by Year in descending order

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Version: 2024-02-01

20
papers

3,281
citations

623734

14
h-index

752698

20
g-index

20
all docs

20
docs citations

20
times ranked

4469
citing authors

#	ARTICLE	IF	CITATIONS
1	Episodes of Diversification and Isolation in Island Southeast Asian and Near Oceanian Male Lineages. <i>Molecular Biology and Evolution</i> , 2022, 39, .	8.9	9
2	Dissecting the paternal founders of Mundari (Austroasiatic) speakers associated with the language dispersal in South Asia. <i>European Journal of Human Genetics</i> , 2021, 29, 528-532.	2.8	2
3	Origin and diffusion of human Y chromosome haplogroup J1-M267. <i>Scientific Reports</i> , 2021, 11, 6659.	3.3	26
4	Phylogenetic history of patrilineages rare in northern and eastern Europe from large-scale re-sequencing of human Y-chromosomes. <i>European Journal of Human Genetics</i> , 2021, 29, 1510-1519.	2.8	1
5	Y-chromosomal connection between Hungarians and geographically distant populations of the Ural Mountain region and West Siberia. <i>Scientific Reports</i> , 2019, 9, 7786.	3.3	12
6	Y-chromosomal analysis of clan structure of Kalmyks, the only European Mongol people, and their relationship to Oirat-Mongols of Inner Asia. <i>European Journal of Human Genetics</i> , 2019, 27, 1466-1474.	2.8	8
7	Investigating the origins of eastern Polynesians using genome-wide data from the Leeward Society Isles. <i>Scientific Reports</i> , 2018, 8, 1823.	3.3	22
8	Genes reveal traces of common recent demographic history for most of the Uralic-speaking populations. <i>Genome Biology</i> , 2018, 19, 139.	8.8	67
9	Origin and spread of human mitochondrial DNA haplogroup U7. <i>Scientific Reports</i> , 2017, 7, 46044.	3.3	25
10	The genetic variation in the R1a clade among the Ashkenazi Levites™ Y chromosome. <i>Scientific Reports</i> , 2017, 7, 14969.	3.3	13
11	“Like sugar in milk” reconstructing the genetic history of the Parsi population. <i>Genome Biology</i> , 2017, 18, 110.	8.8	29
12	Human Y Chromosome Haplogroup N: A Non-trivial Time-Resolved Phylogeography that Cuts across Language Families. <i>American Journal of Human Genetics</i> , 2016, 99, 163-173.	6.2	98
13	Genomic evidence for the Pleistocene and recent population history of Native Americans. <i>Science</i> , 2015, 349, aab3884.	12.6	449
14	A recent bottleneck of Y chromosome diversity coincides with a global change in culture. <i>Genome Research</i> , 2015, 25, 459-466.	5.5	348
15	The genome of a Late Pleistocene human from a Clovis burial site in western Montana. <i>Nature</i> , 2014, 506, 225-229.	27.8	500
16	Unravelling the distinct strains of Tharu ancestry. <i>European Journal of Human Genetics</i> , 2014, 22, 1404-1412.	2.8	17
17	Upper Palaeolithic Siberian genome reveals dual ancestry of Native Americans. <i>Nature</i> , 2014, 505, 87-91.	27.8	821
18	An Aboriginal Australian Genome Reveals Separate Human Dispersals into Asia. <i>Science</i> , 2011, 334, 94-98.	12.6	675

#	ARTICLE	IF	CITATIONS
19	Population Genetic Structure in Indian Austroasiatic Speakers: The Role of Landscape Barriers and Sex-Specific Admixture. <i>Molecular Biology and Evolution</i> , 2011, 28, 1013-1024.	8.9	135
20	Language Shift by Indigenous Population: A Model Genetic Study in South Asia. <i>International Journal of Human Genetics</i> , 2008, 8, 41-50.	0.1	24