Monika Karmin

List of Publications by Year in descending order

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MONIKA KADMIN

#	Article	lF	CITATIONS
1	Upper Palaeolithic Siberian genome reveals dual ancestry of Native Americans. Nature, 2014, 505, 87-91.	27.8	821
2	An Aboriginal Australian Genome Reveals Separate Human Dispersals into Asia. Science, 2011, 334, 94-98.	12.6	675
3	The genome of a Late Pleistocene human from a Clovis burial site in western Montana. Nature, 2014, 506, 225-229.	27.8	500
4	Genomic evidence for the Pleistocene and recent population history of Native Americans. Science, 2015, 349, aab3884.	12.6	449
5	A recent bottleneck of Y chromosome diversity coincides with a global change in culture. Genome Research, 2015, 25, 459-466.	5.5	348
6	Population Genetic Structure in Indian Austroasiatic Speakers: The Role of Landscape Barriers and Sex-Specific Admixture. Molecular Biology and Evolution, 2011, 28, 1013-1024.	8.9	135
7	Human Y Chromosome Haplogroup N: A Non-trivial Time-Resolved Phylogeography that Cuts across Language Families. American Journal of Human Genetics, 2016, 99, 163-173.	6.2	98
8	Genes reveal traces of common recent demographic history for most of the Uralic-speaking populations. Genome Biology, 2018, 19, 139.	8.8	67
9	"Like sugar in milk― reconstructing the genetic history of the Parsi population. Genome Biology, 2017, 18, 110.	8.8	29
10	Origin and diffusion of human Y chromosome haplogroup J1-M267. Scientific Reports, 2021, 11, 6659.	3.3	26
11	Origin and spread of human mitochondrial DNA haplogroup U7. Scientific Reports, 2017, 7, 46044.	3.3	25
12	Language Shift by Indigenous Population: A Model Genetic Study in South Asia. International Journal of Human Genetics, 2008, 8, 41-50.	0.1	24
13	Investigating the origins of eastern Polynesians using genome-wide data from the Leeward Society Isles. Scientific Reports, 2018, 8, 1823.	3.3	22
14	Unravelling the distinct strains of Tharu ancestry. European Journal of Human Genetics, 2014, 22, 1404-1412.	2.8	17
15	The genetic variation in the R1a clade among the Ashkenazi Levites' Y chromosome. Scientific Reports, 2017, 7, 14969.	3.3	13
16	Y-chromosomal connection between Hungarians and geographically distant populations of the Ural Mountain region and West Siberia. Scientific Reports, 2019, 9, 7786.	3.3	12
17	Episodes of Diversification and Isolation in Island Southeast Asian and Near Oceanian Male Lineages. Molecular Biology and Evolution, 2022, 39, .	8.9	9
18	Y-chromosomal analysis of clan structure of Kalmyks, the only European Mongol people, and their relationship to Oirat-Mongols of Inner Asia. European Journal of Human Genetics, 2019, 27, 1466-1474.	2.8	8

#	Article	IF	CITATIONS
19	Dissecting the paternal founders of Mundari (Austroasiatic) speakers associated with the language dispersal in South Asia. European Journal of Human Genetics, 2021, 29, 528-532.	2.8	2
20	Phylogenetic history of patrilineages rare in northern and eastern Europe from large-scale re-sequencing of human X-chromosomes, European Journal of Human Cenetics, 2021, 29, 1510-1519	2.8	1

re-sequencing of human Y-chromosomes. European Journal of Human Genetics, 2021, 29, 1510-1519. 20