

Genomic insights into the formation of human population

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Fine-scale north-to-south genetic admixture profile in Shaanxi Han Chinese revealed by genome-wide demographic history reconstruction. <i>Journal of Systematics and Evolution</i> , 2022, 60, 955-972.	1.6	26
3	Multiple migrations to the Philippines during the last 50,000 years. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	3.3	50
4	Where Asia meets Europe – recent insights from ancient human genomics. <i>Annals of Human Biology</i> , 2021, 48, 191-202.	0.4	5
5	Maternal genetic history of southern East Asians over the past 12,000 years. <i>Journal of Genetics and Genomics</i> , 2021, 48, 899-907.	1.7	6
6	Significant East Asian Affinity of the Sichuan Hui Genomic Structure Suggests the Predominance of the Cultural Diffusion Model in the Genetic Formation Process. <i>Frontiers in Genetics</i> , 2021, 12, 626710.	1.1	17
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10	Genomic Insights Into the Admixture History of Mongolic- and Tungusic-Speaking Populations From Southwestern East Asia. <i>Frontiers in Genetics</i> , 2021, 12, 685285.	1.1	23
12	Genome Analysis Traces Regional Dispersal of Rice in Taiwan and Southeast Asia. <i>Molecular Biology and Evolution</i> , 2021, 38, 4832-4846.	3.5	16
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19	Philippine Ayta possess the highest level of Denisovan ancestry in the world. <i>Current Biology</i> , 2021, 31, 4219-4230.e10.	1.8	37
20	Genome of a middle Holocene hunter-gatherer from Wallacea. <i>Nature</i> , 2021, 596, 543-547.	13.7	35
21	Ancient Mitochondrial Genomes Reveal Extensive Genetic Influence of the Steppe Pastoralists in Western Xinjiang. <i>Frontiers in Genetics</i> , 2021, 12, 740167.	1.1	6
22	Genetic insights into the paternal admixture history of Chinese Mongolians via high-resolution customized Y-SNP SNaPshot panels. <i>Forensic Science International: Genetics</i> , 2021, 54, 102565.	1.6	19
23	Dairying enabled Early Bronze Age Yamnaya steppe expansions. <i>Nature</i> , 2021, 598, 629-633.	13.7	47

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24	Insights into human history from the first decade of ancient human genomics. <i>Science</i> , 2021, 373, 1479-1484.	6.0	41
25	Genomic Insight Into the Population Structure and Admixture History of Tai-Kadai-Speaking Sui People in Southwest China. <i>Frontiers in Genetics</i> , 2021, 12, 735084.	1.1	13
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36	Female craniometrics support the â€™two-layer modelâ€™ of human dispersal in Eastern Eurasia. <i>Scientific Reports</i> , 2021, 11, 20830.	1.6	7
37	The genomic origins of the Bronze Age Tarim Basin mummies. <i>Nature</i> , 2021, 599, 256-261.	13.7	65
39	Genomic history and forensic characteristics of Sherpa highlanders on the Tibetan Plateau inferred from high-resolution InDel panel and genome-wide SNPs. <i>Forensic Science International: Genetics</i> , 2022, 56, 102633.	1.6	13
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44	Genomic insights into Neolithic farming-related migrations in the junction of east and southeast Asia. <i>American Journal of Biological Anthropology</i> , 2022, 177, 328-342.	0.6	11
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80	Genomic analyses of 10,376 individuals in the Westlake BioBank for Chinese (WBBC) pilot project. <i>Nature Communications</i> , 2022, 13, .	5.8	41
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91	Genomic Insights Into the Unique Demographic History and Genetic Structure of Five Hmong-Mien-Speaking Miao and Yao Populations in Southwest China. <i>Frontiers in Ecology and Evolution</i> , 0, 10, .	1.1	8
92	Northeastern Asian and Jomon-related genetic structure in the Three Kingdoms period of Gimhae, Korea. <i>Current Biology</i> , 2022, 32, 3232-3244.e6.	1.8	6
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