

Multiple Deeply Divergent Denisovan Ancestries in Pap

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Aspects of human physical and behavioural evolution during the last 1 million years. <i>Journal of Quaternary Science</i> , 2019, 34, 355-378.	1.1	63
2	The Northern Route for Human dispersal in Central and Northeast Asia: New evidence from the site of Tolbor-16, Mongolia. <i>Scientific Reports</i> , 2019, 9, 11759.	1.6	55
3	Using hominin introgression to trace modern human dispersals. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 15327-15332.	3.3	23
4	Genetics, adaptation to environmental changes and archaic admixture in the pathogenesis of diabetes mellitus in Indigenous Australians. <i>Reviews in Endocrine and Metabolic Disorders</i> , 2019, 20, 321-332.	2.6	2
6	Sex-linked genetic diversity originates from persistent sociocultural processes at microgeographic scales. <i>Royal Society Open Science</i> , 2019, 6, 190733.	1.1	0
7	The first hominin fleet. <i>Nature Ecology and Evolution</i> , 2019, 3, 999-1000.	3.4	10
8	Searching for archaic contribution in Africa. <i>Annals of Human Biology</i> , 2019, 46, 129-139.	0.4	4
9	Experimental futures in archaeology. <i>Antiquity</i> , 2019, 93, 808-810.	0.5	5
10	Archaic mitochondrial DNA inserts in modern day nuclear genomes. <i>BMC Genomics</i> , 2019, 20, 1017.	1.2	8
11	Archaic hominin introgression into modern human genomes. <i>American Journal of Physical Anthropology</i> , 2020, 171, 60-73.	2.1	33
12	New portraits of the Denisovans. <i>Science Bulletin</i> , 2020, 65, 1-3.	4.3	4
13	Last appearance of <i>Homo erectus</i> at Ngandong, Java, 117,000–108,000 years ago. <i>Nature</i> , 2020, 577, 381-385.	13.7	97
14	The Impact of Ancient Genome Studies in Archaeology. <i>Annual Review of Anthropology</i> , 2020, 49, 277-298.	0.4	13
15	Human evolutionary history in Eastern Eurasia using insights from ancient DNA. <i>Current Opinion in Genetics and Development</i> , 2020, 62, 78-84.	1.5	18
16	Global Picture of Genetic Relatedness and the Evolution of Humankind. <i>Biology</i> , 2020, 9, 392.	1.3	2
17	HIV-1 p24Gag adaptation to modern and archaic HLA-allele frequency differences in ethnic groups contributes to viral subtype diversification. <i>Virus Evolution</i> , 2020, 6, veaa085.	2.2	7
18	Mapping gene flow between ancient hominins through demography-aware inference of the ancestral recombination graph. <i>PLoS Genetics</i> , 2020, 16, e1008895.	1.5	76
19	Methods for detecting introgressed archaic sequences. <i>Current Opinion in Genetics and Development</i> , 2020, 62, 85-90.	1.5	6

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21	The Spatial Signature of Introgression After a Biological Invasion With Hybridization. <i>Frontiers in Ecology and Evolution</i> , 2020, 8, .	1.1	20
22	A different paradigm for the colonisation of Sahul. <i>Archaeology in Oceania</i> , 2020, 55, 182-191.	0.3	1
23	Two-stage mid-Brunhes climate transition and mid-Pleistocene human diversification. <i>Earth-Science Reviews</i> , 2020, 210, 103354.	4.0	35
24	The reversal of human phylogeny: Homo left Africa as erectus, came back as sapiens sapiens. <i>Hereditas</i> , 2020, 157, 51.	0.5	3
25	A Revised Model of Anatomically Modern Human Expansions Out of Africa through a Machine Learning Approximate Bayesian Computation Approach. <i>Genes</i> , 2020, 11, 1510.	1.0	4
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35	Neanderthal-Denisovan ancestors interbred with a distantly related hominin. <i>Science Advances</i> , 2020, 6, eaay5483.	4.7	43
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64	Phenotypic differences between highlanders and lowlanders in Papua New Guinea. <i>PLoS ONE</i> , 2021, 16, e0253921.	1.1	4
65	Papua New Guinean Genomes Reveal the Complex Settlement of North Sahul. <i>Molecular Biology and Evolution</i> , 2021, 38, 5107-5121.	3.5	11
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