

Complex Patterns of Admixture across the Indonesian A

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Kinship structures create persistent channels for language transmission. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 12910-12915.	7.1	18
2	Genetic relatedness of indigenous ethnic groups in northern Borneo to neighboring populations from Southeast Asia, as inferred from genome-wide SNP data. Annals of Human Genetics, 2018, 82, 216-226.	0.8	13
3	Investigating the origins of eastern Polynesians using genome-wide data from the Leeward Society Isles. Scientific Reports, 2018, 8, 1823.	3.3	22
4	The Gateway from Near into Remote Oceania: New Insights from Genome-Wide Data. Molecular Biology and Evolution, 2018, 35, 871-886.	8.9	38
5	Genes reveal traces of common recent demographic history for most of the Uralic-speaking populations. Genome Biology, 2018, 19, 139.	8.8	67
6	The prehistoric peopling of Southeast Asia. Science, 2018, 361, 88-92.	12.6	291
7	Big genetic data and its big data protection challenges. Computer Law and Security Review, 2018, 34, 1000-1018.	2.2	20
8	Somewhere beyond the sea: Human cranial remains from the Lesser Sunda Islands (Alor Island.) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Evolution, 2019, 134, 102638.	2.6	13
9	Using hominin introgression to trace modern human dispersals. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 15327-15332.	7.1	23
10	Uganda Genome Resource Enables Insights into Population History and Genomic Discovery in Africa. Cell, 2019, 179, 984-1002.e36.	28.9	152
11	Sex-linked genetic diversity originates from persistent sociocultural processes at microgeographic scales. Royal Society Open Science, 2019, 6, 190733.	2.4	0
12	The Importance of Ethnicity Definitions and Pharmacogenomics in Ethnobridging and Pharmacovigilance. , 2019, , 289-327.		1
13	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
14	Population genetics, diversity and forensic characteristics of Tai-Kadai-speaking Bouyei revealed by insertion/deletions markers. Molecular Genetics and Genomics, 2019, 294, 1343-1357.	2.1	32
15	The dispersal of Austronesian languages in Island South East Asia: Current findings and debates. Language and Linguistics Compass, 2019, 13, e12325.	2.3	13
16	Multiple Deeply Divergent Denisovan Ancestries in Papuans. Cell, 2019, 177, 1010-1021.e32.	28.9	181
17	Denisovan, modern human and mouse TNFAIP3 alleles tune A20 phosphorylation and immunity. Nature Immunology, 2019, 20, 1299-1310.	14.5	53
18	Indonesians Human Leukocyte Antigen (HLA) Distributions and Correlations with Global Diseases. Immunological Investigations, 2020, 49, 333-363.	2.0	8

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19	The paternal and maternal genetic history of Vietnamese populations. <i>European Journal of Human Genetics</i> , 2020, 28, 636-645.	2.8	24
20	A Likelihood Approach for Uncovering Selective Sweep Signatures from Haplotype Data. <i>Molecular Biology and Evolution</i> , 2020, 37, 3023-3046.	8.9	28
21	Mitogenomics of macaques (<i>Macaca</i>) across Wallace's Line in the context of modern human dispersals. <i>Journal of Human Evolution</i> , 2020, 146, 102852.	2.6	18
22	Differences in local population history at the finest level: the case of the Estonian population. <i>European Journal of Human Genetics</i> , 2020, 28, 1580-1591.	2.8	23
23	A different paradigm for the colonisation of Sahul. <i>Archaeology in Oceania</i> , 2020, 55, 182-191.	0.7	1
24	Genome-wide DNA methylation and gene expression patterns reflect genetic ancestry and environmental differences across the Indonesian archipelago. <i>PLoS Genetics</i> , 2020, 16, e1008749.	3.5	30
25	Ritual tooth ablation and the Austronesian expansion: Evidence from eastern Indonesia and the Pacific Islands. <i>Journal of Island and Coastal Archaeology</i> , 2020, , 1-32.	1.4	3
26	A different paradigm for the initial colonisation of Sahul. <i>Archaeology in Oceania</i> , 2020, 55, 1-14.	0.7	13
27	Genomic landscape of the signals of positive natural selection in populations of Northern Eurasia: A view from Northern Russia. <i>PLoS ONE</i> , 2020, 15, e0228778.	2.5	6
28	Artificial intelligence-assisted prediction of preeclampsia: Development and external validation of a nationwide health insurance dataset of the BPJS Kesehatan in Indonesia. <i>EBioMedicine</i> , 2020, 54, 102710.	6.1	33
30	Human genetic admixture. <i>PLoS Genetics</i> , 2021, 17, e1009374.	3.5	28
31	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, .	7.1	50
32	A Mix of Old British and Modern European Breeds: Genomic Prediction of Breed Composition of Smallholder Pigs in Uganda. <i>Frontiers in Genetics</i> , 2021, 12, 676047.	2.3	7
33	Mitogenomes Reveal Two Major Influxes of Papuan Ancestry across Wallacea Following the Last Glacial Maximum and Austronesian Contact. <i>Genes</i> , 2021, 12, 965.	2.4	15
34	Papua New Guinean Genomes Reveal the Complex Settlement of North Sahul. <i>Molecular Biology and Evolution</i> , 2021, 38, 5107-5121.	8.9	11
35	Ancient DNA from Guam and the peopling of the Pacific. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	25
45	Reconstruction of the Austronesian Diaspora in the Era of Genomics. <i>Human Biology</i> , 2020, 92, 247.	0.2	6
48	Genetic architecture of gene regulation in Indonesian populations identifies QTLs associated with global and local ancestries. <i>American Journal of Human Genetics</i> , 2022, 109, 50-65.	6.2	11

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49	The Peopling and Migration History of the Natives in Peninsular Malaysia and Borneo: A Glimpse on the Studies Over the Past 100 years. <i>Frontiers in Genetics</i> , 2022, 13, 767018.	2.3	1
50	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
52	Ancient genomes from the last three millennia support multiple human dispersals into Wallacea. <i>Nature Ecology and Evolution</i> , 2022, 6, 1024-1034.	7.8	15
53	Chronology of natural selection in Oceanian genomes. <i>IScience</i> , 2022, 25, 104583.	4.1	3
54	Sequence analyses of Malaysian Indigenous communities reveal historical admixture between Hoabinhian hunter-gatherers and Neolithic farmers. <i>Scientific Reports</i> , 2022, 12, .	3.3	1
55	Talking Dead. New burials from Tron Bon Lei (Alor Island, Indonesia) inform on the evolution of mortuary practices from the terminal Pleistocene to the Holocene in Southeast Asia. <i>PLoS ONE</i> , 2022, 17, e0267635.	2.5	6
56	Mainland-coastal interactions in East Borneo: Inter-site comparison and Bayesian chronological models of two Late Pleistoceneâ€“Holocene sequences (Liang Abu and Kimanis rock shelters). <i>Journal of Island and Coastal Archaeology</i> , 0, , 1-38.	1.4	1
58	A Cloth that Promises Resurrection. <i>Indonesia and the Malay World</i> , 0, , 1-29.	0.9	0
59	The loss of biodiversity in Madagascar is contemporaneous with major demographic events. <i>Current Biology</i> , 2022, , .	3.9	1
60	Human Genetic Research in Wallacea and Sahul: Recent Findings and Future Prospects. <i>Genes</i> , 2022, 13, 2373.	2.4	1
61	Denisovan introgression has shaped the immune system of present-day Papuans. <i>PLoS Genetics</i> , 2022, 18, e1010470.	3.5	9
63	Prehistoric Human Migrations in Southeast Asia through the Lenses of Burial Practices. , 0, , .		0
64	Î² Thalassemia Mutation Flow in Indonesia: A Migration Perspective. <i>Thalassemia Reports</i> , 2023, 13, 253-261.	0.5	0
65	The Effect of Cognitive Load on Learning Memory of Online Learning Accounting Students in the Philippines. <i>Sustainability</i> , 2024, 16, 1686.	3.2	0