A Neolithic expansion, but strong genetic structure, in t Guinea

Science 357, 1160-1163

DOI: 10.1126/science.aan3842

Citation Report

#	Article	IF	CITATIONS
1	The Gateway from Near into Remote Oceania: New Insights from Genome-Wide Data. Molecular Biology and Evolution, 2018, 35, 871-886.	8.9	38
2	Whole-genome sequencing of 175 Mongolians uncovers population-specific genetic architecture and gene flow throughout North and East Asia. Nature Genetics, 2018, 50, 1696-1704.	21.4	38
4	Human Genetic Variation and HIV/AIDS in Papua New Guinea: Time to Connect the Dots. Current HIV/AIDS Reports, 2018, 15, 431-440.	3.1	2
5	The evolutionary history and human settlement of Australia and the Pacific. Current Opinion in Genetics and Development, 2018, 53, 53-59.	3.3	11
6	Adaptive archaic introgression of copy number variants and the discovery of previously unknown human genes. Science, 2019, 366, .	12.6	65
7	Multiple Deeply Divergent Denisovan Ancestries in Papuans. Cell, 2019, 177, 1010-1021.e32.	28.9	181
8	Giving it a burl: towards the integration of genetics, isotope chemistry, and osteoarchaeology in Cape York, Tropical North Queensland, Australia. World Archaeology, 2019, 51, 602-619.	1.1	20
9	Contributions of Quaternary botany to modern ecology and biogeography. Plant Ecology and Diversity, 2019, 12, 189-385.	2.4	103
10	Weighing outcome vs. intent across societies: How cultural models of mind shape moral reasoning. Cognition, 2019, 182, 95-108.	2.2	67
11	A different paradigm for the colonisation of Sahul. Archaeology in Oceania, 2020, 55, 182-191.	0.7	1
12	Papuan mitochondrial genomes and the settlement of Sahul. Journal of Human Genetics, 2020, 65, 875-887.	2.3	24
13	Emergence of a Neolithic in highland New Guinea by 5000 to 4000 years ago. Science Advances, 2020, 6, eaay4573.	10.3	18
14	Insights into human genetic variation and population history from 929 diverse genomes. Science, 2020, 367, .	12.6	534
15	Small game hunting in montane rainforests: Specialised capture and broad spectrum foraging in the Late Pleistocene to Holocene New Guinea Highlands. Quaternary Science Reviews, 2021, 253, 106742.	3.0	11
16	Genomic insights into population history and biological adaptation in Oceania. Nature, 2021, 592, 583-589.	27.8	100
17	Mitogenomes Reveal Two Major Influxes of Papuan Ancestry across Wallacea Following the Last Glacial Maximum and Austronesian Contact. Genes, 2021, 12, 965.	2.4	15
18	Phenotypic differences between highlanders and lowlanders in Papua New Guinea. PLoS ONE, 2021, 16, e0253921.	2.5	4
19	Papua New Guinean Genomes Reveal the Complex Settlement of North Sahul. Molecular Biology and Evolution, 2021, 38, 5107-5121.	8.9	11

#	Article	IF	CITATIONS
20	Philippine Ayta possess the highest level of Denisovan ancestry in the world. Current Biology, 2021, 31, 4219-4230.e10.	3.9	37
21	Late Pleistocene/Early Holocene sites in the montane forests of New Guinea yield early record of cassowary hunting and egg harvesting. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	8
22	A contextualised review of genomic evidence for gene flow events between Papuans and Indigenous Australians in Cape York, Queensland. Quaternary International, 2021, 603, 22-30.	1.5	6
23	Ancient DNA from Guam and the peopling of the Pacific. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	25
25	Parkinsonism and motor neuron disorders: Lessons from Western Pacific ALS/PDC. Journal of the Neurological Sciences, 2022, 433, 120021.	0.6	12
26	Reconsidering the †Neolithic' at Manim rock shelter, Wurup Valley, Papua New Guinea. , 2019, , 81-99.		2
27	Reconstruction of the Austronesian Diaspora in the Era of Genomics. Human Biology, 2020, 92, 247.	0.2	6
29	A dentateâ€stamped Lapita dish from the central south coast of Papua. Archaeology in Oceania, 0, , .	0.7	0
31	Episodes of Diversification and Isolation in Island Southeast Asian and Near Oceanian Male Lineages. Molecular Biology and Evolution, 2022, 39, .	8.9	9
32	Late Quaternary changes in malaria-free areas in Papua New Guinea and the future perspectives. Quaternary International, 2022, 628, 28-43.	1.5	2
33	The Tibetan-Yi region is both a corridor and a barrier for human gene flow. Cell Reports, 2022, 39, 110720.	6.4	8
34	Prisoners of a distant past? Linguistic diversity and the time-depth of human settlement in Papua New Guinea. World Development, 2022, 157, 105921.	4.9	0
35	Ancient DNA reveals five streams of migration into Micronesia and matrilocality in early Pacific seafarers. Science, 2022, 377, 72-79.	12.6	13
36	Chronology of natural selection in Oceanian genomes. IScience, 2022, 25, 104583.	4.1	3
37	Assessing Human Genome-wide Variation in the Massim Region of Papua New Guinea and Implications for the Kula Trading Tradition. Molecular Biology and Evolution, 2022, 39, .	8.9	0
39	Genomic perspectives on human dispersals during the Holocene. Proceedings of the National Academy of Sciences of the United States of America, 2023, 120, .	7.1	12
40	Is there still evolution in the human population?. Biologia Futura, 2022, 73, 359-374.	1.4	1
41	Fit for purpose: investigating adaptations in late Pleistocene lithic technology to an island environment at Buang Merabak, New Ireland, Papua New Guinea. World Archaeology, 2022, 54, 317-337.	1.1	0

#	Article	IF	CITATIONS
42	Testing of two SNP array–based genealogy algorithms using extended Han Chinese pedigrees and recommendations for improved performances in forensic practice. Electrophoresis, 0, , .	2.4	0
43	Insights from ancient human DNA into the colonization of Oceania. , 2023, , .		0
44	Indigenous Australian genomes show deep structure and rich novel variation. Nature, 0, , .	27.8	1
45	Population structure and migration in the Eastern Highlands of Papua New Guinea, a region impacted by the kuru epidemic. American Journal of Human Genetics, 2024, 111, 668-679.	6.2	0