## **Reconstructing Prehistoric African Population Structur**

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Citation Report

#	Article	IF	CITATIONS
1	Later Stone Age toolstone acquisition in the Central Rift Valley of Kenya: Portable XRF of Eburran obsidian artifacts from Leakey's excavations at Gamble's Cave II. Journal of Archaeological Science: Reports, 2018, 18, 475-486.	0.2	8
2	Insights into Modern Human Prehistory Using Ancient Genomes. Trends in Genetics, 2018, 34, 184-196.	2.9	50
3	Mediterranean Y-chromosome 2.0—why the Y in the Mediterranean is still relevant in the postgenomic era. Annals of Human Biology, 2018, 45, 20-33.	0.4	8
4	How have our clocks evolved? Adaptive and demographic history of the out-of-African dispersal told by polymorphic loci in circadian genes. Chronobiology International, 2018, 35, 511-532.	0.9	7
5	Drivers and trajectories of land cover change in East Africa: Human and environmental interactions from 6000†years ago to present. Earth-Science Reviews, 2018, 178, 322-378.	4.0	129
6	Tales of Human Migration, Admixture, and Selection in Africa. Annual Review of Genomics and Human Genetics, 2018, 19, 405-428.	2.5	78
7	Ancient Genomics of Modern Humans: The First Decade. Annual Review of Genomics and Human Genetics, 2018, 19, 381-404.	2.5	161
8	Pleistocene North African genomes link Near Eastern and sub-Saharan African human populations. Science, 2018, 360, 548-552.	6.0	142
9	The demographic history and mutational load of African hunter-gatherers and farmers. Nature Ecology and Evolution, 2018, 2, 721-730.	3.4	38
10	Ecological risk, demography and technological complexity in the Late Pleistocene of northern Malawi: implications for geographical patterning in the Middle Stone Age. Journal of Quaternary Science, 2018, 33, 261-284.	1.1	21
11	Recent advances in understanding evolution of the placenta: insights from transcriptomics. F1000Research, 2018, 7, 89.	0.8	12
12	Clarifying distinct models of modern human origins in Africa. Current Opinion in Genetics and Development, 2018, 53, 148-156.	1.5	51
13	Ancient human DNA: How sequencing the genome of a boy from Ballito Bay changed human history. South African Journal of Science, 2018, 114, 3.	0.3	5
14	Principles of Archaeogenetics and the Current Trends of Ancient Genome Studies. Korean Journal of Physical Anthropology, 2018, 31, 105.	0.2	0
15	Population Genetic Diversity and Phylogenetic Characteristics for High-Altitude Adaptive Kham Tibetan Revealed by DNATyperTM 19 Amplification System. Frontiers in Genetics, 2018, 9, 630.	1.1	17
16	The evolutionary history of Southern Africa. Current Opinion in Genetics and Development, 2018, 53, 157-164.	1.5	10
17	Quantifying and reducing spurious alignments for the analysis of ultra-short ancient DNA sequences. BMC Biology, 2018, 16, 121.	1.7	41
18	Human evolutionary history of Eastern Africa. Current Opinion in Genetics and Development, 2018, 53, 134-139.	1.5	8

#	Article	IF	CITATIONS
19	Rare genetic sequences illuminate early humans' history in Africa. Nature, 2018, 563, 13-14.	13.7	0
20	Genetic structure and sexâ€biased gene flow in the history of southern African populations. American Journal of Physical Anthropology, 2018, 167, 656-671.	2.1	25
21	The Year In Genetic Anthropology: New Lands, New Technologies, New Questions. American Anthropologist, 2018, 120, 266-277.	0.7	5
22	Boots on the ground in Africa's ancient DNA â€~revolution': archaeological perspectives on ethics and best practices. Antiquity, 2018, 92, 803-815.	0.5	63
23	Middle Stone Age human teeth from Magubike rockshelter, Iringa Region, Tanzania. PLoS ONE, 2018, 13, e0200530.	1.1	14
24	Biological Sexing of a 4000-Year-Old Egyptian Mummy Head to Assess the Potential of Nuclear DNA Recovery from the Most Damaged and Limited Forensic Specimens. Genes, 2018, 9, 135.	1.0	39
25	Did Our Species Evolve in Subdivided Populations across Africa, and Why Does It Matter?. Trends in Ecology and Evolution, 2018, 33, 582-594.	4.2	315
26	Crafting Early African Histories with Jan Vansina. History in Africa, 2018, 45, 99-112.	0.1	1
27	A history of male migration in and out of the Green Sahara. Genome Biology, 2018, 19, 30.	3.8	2
28	African genetic diversity provides novel insights into evolutionary history and local adaptations. Human Molecular Genetics, 2018, 27, R209-R218.	1.4	38
29	Archaeogenomic evidence from the southwestern US points to a pre-Hispanic scarlet macaw breeding colony. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 8740-8745.	3.3	20
30	Ancient DNA from Chalcolithic Israel reveals the role of population mixture in cultural transformation. Nature Communications, 2018, 9, 3336.	5.8	71
31	Rivals No More: Jan Vansina, Precolonial African Historiography, and Archaeology. History in Africa, 2018, 45, 145-160.	0.1	3
32	Becoming Neolithic in words, thoughts and deeds. Journal of Social Archaeology, 2019, 19, 67-91.	1.0	6
33	Ancient DNA from the skeletons of Roopkund Lake reveals Mediterranean migrants in India. Nature Communications, 2019, 10, 3670.	5.8	19
34	Direct Evidence of an Increasing Mutational Load in Humans. Molecular Biology and Evolution, 2019, 36, 2823-2829.	3.5	12
35	Identifying loci under positive selection in complex population histories. Genome Research, 2019, 29, 1506-1520.	2.4	36
36	Genetic Affinities among Southern Africa Hunter-Gatherers and the Impact of Admixing Farmer and Herder Populations. Molecular Biology and Evolution, 2019, 36, 1849-1861.	3.5	21

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#	Article	IF	CITATIONS
37	Population structure and the evolution of <i>Homo sapiens</i> in Africa. Evolutionary Anthropology, 2019, 28, 179-188.	1.7	25
38	Diet and adult ageâ€atâ€death among mobile foragers: A synthesis of bioarcheological methods. American Journal of Physical Anthropology, 2019, 170, 131-147.	2.1	9
39	The long walk to African genomics. Genome Biology, 2019, 20, 130.	3.8	27
40	Unraveling African diversity from a crossâ€disciplinary perspective. Evolutionary Anthropology, 2019, 28, 288-292.	1.7	1
41	Uganda Genome Resource Enables Insights into Population History and Genomic Discovery in Africa. Cell, 2019, 179, 984-1002.e36.	13.5	152
42	Integrating evidence of land use and land cover change for land management policy formulation along the Kenya-Tanzania borderlands. Anthropocene, 2019, 28, 100228.	1.6	10
43	Population Structure and Implications on the Genetic Architecture of HIV-1 Phenotypes Within Southern Africa. Frontiers in Genetics, 2019, 10, 905.	1.1	15
44	Balancing analytical goals and anthropological stewardship in the midst of the paleogenomics revolution. World Archaeology, 2019, 51, 560-573.	0.5	17
45	Out of Africa by spontaneous migration waves. PLoS ONE, 2019, 14, e0201998.	1.1	15
46	Genomics of disease risk in globally diverse populations. Nature Reviews Genetics, 2019, 20, 520-535.	7.7	217
47	Ancient DNA reveals a multistep spread of the first herders into sub-Saharan Africa. Science, 2019, 365,	6.0	96
48	Models of archaic admixture and recent history from two-locus statistics. PLoS Genetics, 2019, 15, e1008204.	1.5	57
49	A catalog of single nucleotide changes distinguishing modern humans from archaic hominins. Scientific Reports, 2019, 9, 8463.	1.6	60
50	A Rare Deep-Rooting D0 African Y-Chromosomal Haplogroup and Its Implications for the Expansion of Modern Humans Out of Africa. Genetics, 2019, 212, 1421-1428.	1.2	35
51	PopNetD3—A Network-Based Web Resource for Exploring Population Structure. Genome Biology and Evolution, 2019, 11, 1730-1735.	1.1	1
52	The lithic assemblage from Sugenya, a Pastoral Neolithic site of the Elmenteitan tradition in southwestern Kenya. Azania, 2019, 54, 4-32.	0.4	2
53	How well do we understand the basis of classic selective sweeps in humans?. FEBS Letters, 2019, 593, 1431-1448.	1.3	17
54	Assessing the importance of cultural diffusion in the Bantu spread into southeastern Africa. PLoS ONE, 2019, 14, e0215573.	1.1	13

#	Article	IF	CITATIONS
55	Population genetic structure in Fennoscandian landrace rye (Secale cereale L.) spanning 350Âyears. Genetic Resources and Crop Evolution, 2019, 66, 1059-1071.	0.8	6
56	African evolutionary history inferred from whole genome sequence data of 44 indigenous African populations. Genome Biology, 2019, 20, 82.	3.8	84
57	Whole-genome sequence analysis of a Pan African set of samples reveals archaic gene flow from an extinct basal population of modern humans into sub-Saharan populations. Genome Biology, 2019, 20, 77.	3.8	50
58	A dispersal of Homo sapiens from southern to eastern Africa immediately preceded the out-of-Africa migration. Scientific Reports, 2019, 9, 4728.	1.6	49
59	Analyses of Neanderthal introgression suggest that Levantine and southern Arabian populations have a shared population history. American Journal of Physical Anthropology, 2019, 169, 227-239.	2.1	11
60	Isolating the human cochlea to generate bone powder for ancient DNA analysis. Nature Protocols, 2019, 14, 1194-1205.	5.5	54
61	Ancestral mitochondrial N lineage from the Neolithic â€~green' Sahara. Scientific Reports, 2019, 9, 3530.	1.6	10
62	A western route of prehistoric human migration from Africa into the Iberian Peninsula. Proceedings of the Royal Society B: Biological Sciences, 2019, 286, 20182288.	1.2	47
63	Enamel pearls: Their occurrence in recent human populations and earliest manifestation in the modern human lineage. Archives of Oral Biology, 2019, 101, 147-155.	0.8	2
64	Human Immunology through the Lens of Evolutionary Genetics. Cell, 2019, 177, 184-199.	13.5	105
65	Genome-wide data from the Bubi of Bioko Island clarifies the Atlantic fringe of the Bantu dispersal. BMC Genomics, 2019, 20, 179.	1.2	6
66	The genomic history of the Iberian Peninsula over the past 8000 years. Science, 2019, 363, 1230-1234.	6.0	340
67	The Emergence of Genomic Research in Africa and New Frameworks for Equity in Biomedical Research. Ethnicity and Disease, 2019, 29, 179-186.	1.0	18
68	Positive selection in Europeans and East-Asians at the ABCA12 gene. Scientific Reports, 2019, 9, 4843.	1.6	1
69	Genome-Wide Characterization of Arabian Peninsula Populations: Shedding Light on the History of a Fundamental Bridge between Continents. Molecular Biology and Evolution, 2019, 36, 575-586.	3.5	45
70	Ancient DNA Analysis of the Thulamela Remains: Deciphering the Migratory Patterns of a Southern African Population. Journal of African Archaeology, 2019, 17, 161-172.	0.3	2
71	An ancient DNA Pacific journey: a case study of collaboration between archaeologists and geneticists. World Archaeology, 2019, 51, 620-639.	0.5	7
72	Ancient DNA. , 2019, , 13-34.		0

#	Article	IF	CITATIONS
73	Unlocking the origins and biology of domestic animals using ancient DNA and paleogenomics. BMC Biology, 2019, 17, 98.	1.7	48
74	Ostrich eggshell bead diameter in the Holocene: Regional variation with the spread of herding in eastern and southern Africa. PLoS ONE, 2019, 14, e0225143.	1.1	12
75	West Asian sources of the Eurasian component in Ethiopians: a reassessment. Scientific Reports, 2019, 9, 18811.	1.6	14
76	HOPS: automated detection and authentication of pathogen DNA in archaeological remains. Genome Biology, 2019, 20, 280.	3.8	67
77	Human origins in a southern African palaeo-wetland and first migrations. Nature, 2019, 575, 185-189.	13.7	79
78	Long-term Trends in Terrestrial and Marine Invertebrate Exploitation on the Eastern African Coast: Insights from Kuumbi Cave, Zanzibar. Journal of Island and Coastal Archaeology, 2019, 14, 479-514.	0.6	9
79	The people behind the samples: Biographical features of Past Hunter-Gatherers from KwaZulu-Natal who yielded aDNA. International Journal of Paleopathology, 2019, 24, 158-164.	0.8	11
80	Human teeth from securely stratified Middle Stone Age contexts at Sibudu, South Africa. Archaeological and Anthropological Sciences, 2019, 11, 3491-3501.	0.7	5
81	Approximate Bayesian computation with deep learning supports a third archaic introgression in Asia and Oceania. Nature Communications, 2019, 10, 246.	5.8	97
82	Limits of long-term selection against Neandertal introgression. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 1639-1644.	3.3	151
83	Sorting the sheep from the goats in the Pastoral Neolithic: morphological and biomolecular approaches at Luxmanda, Tanzania. Archaeological and Anthropological Sciences, 2019, 11, 3047-3062.	0.7	28
84	Off-target phenotypes in forensic DNA phenotyping and biogeographic ancestry inference: A resource. Forensic Science International: Genetics, 2019, 38, 93-104.	1.6	15
85	Along the Indian Ocean Coast: Genomic Variation in Mozambique Provides New Insights into the Bantu Expansion. Molecular Biology and Evolution, 2020, 37, 406-416.	3.5	32
86	Archaic hominin introgression into modern human genomes. American Journal of Physical Anthropology, 2020, 171, 60-73.	2.1	33
87	Late Pleistocene to Holocene human palaeoecology in the tropical environments of coastal eastern Africa. Palaeogeography, Palaeoclimatology, Palaeoecology, 2020, 537, 109438.	1.0	37
88	A different view on fine-scale population structure in Western African populations. Human Genetics, 2020, 139, 45-59.	1.8	13
89	The Impact of Ancient Genome Studies in Archaeology. Annual Review of Anthropology, 2020, 49, 277-298.	0.4	13
90	Y-Chromosome Variation in Southern African Khoe-San Populations Based on Whole-Genome Sequences. Genome Biology and Evolution, 2020, 12, 1031-1039.	1.1	6

#	Article	IF	CITATIONS
91	Khoe-San Genomes Reveal Unique Variation and Confirm the Deepest Population Divergence in Homo sapiens. Molecular Biology and Evolution, 2020, 37, 2944-2954.	3.5	60
92	Middle Stone Age lithic assemblages from Leba Cave (Southwest Angola). Journal of Archaeological Science: Reports, 2020, 32, 102413.	0.2	4
93	A Late Pleistocene human humerus from Rusinga Island, Lake Victoria, Kenya. Journal of Human Evolution, 2020, 146, 102855.	1.3	5
94	Methods for detecting introgressed archaic sequences. Current Opinion in Genetics and Development, 2020, 62, 85-90.	1.5	6
95	Intra-tooth stable isotope profiles in warthog canines and third molars: Implications for paleoenvironmental reconstructions. Chemical Geology, 2020, 554, 119799.	1.4	6
96	Introduction to special issue: Frontiers in baboon research. Journal of Human Evolution, 2020, 146, 102822.	1.3	5
97	Human adaptation over the past 40,000 years. Current Opinion in Genetics and Development, 2020, 62, 97-104.	1.5	20
98	Current Trends in Ancient DNA Study. , 2020, , 1-16.		Ο
99	ContamLD: estimation of ancient nuclear DNA contamination using breakdown of linkage disequilibrium. Genome Biology, 2020, 21, 199.	3.8	29
100	High-depth African genomes inform human migration and health. Nature, 2020, 586, 741-748.	13.7	197
101	Origins and genetic legacy of prehistoric dogs. Science, 2020, 370, 557-564.	6.0	152
103	Evolution of Human Brain Size-Associated NOTCH2NL Genes Proceeds toward Reduced Protein Levels. Molecular Biology and Evolution, 2020, 37, 2531-2548.	3.5	10
104	Ancient genomes reveal complex patterns of population movement, interaction, and replacement in sub-Saharan Africa. Science Advances, 2020, 6, eaaz0183.	4.7	56
105	African population history: an ancient DNA perspective. Current Opinion in Genetics and Development, 2020, 62, 8-15.	1.5	29
106	Trajectories of cultural innovation from the Middle to Later Stone Age in Eastern Africa: Personal ornaments, bone artifacts, and ocher from Panga ya Saidi, Kenya. Journal of Human Evolution, 2020, 141, 102737.	1.3	47
107	Tracking human population structure through time from whole genome sequences. PLoS Genetics, 2020, 16, e1008552.	1.5	71
108	Insights into human genetic variation and population history from 929 diverse genomes. Science, 2020, 367, .	6.0	534
109	An isotopic test of the seasonal migration hypothesis for large grazing ungulates inhabiting the Palaeo-Agulhas Plain. Quaternary Science Reviews, 2020, 235, 106221.	1.4	10

#	Article	IF	CITATIONS
110	A southern African archaeological database of organic containers and materials, 800 cal BC to cal AD 1500: Possible implications for the transition from foraging to livestock-keeping. PLoS ONE, 2020, 15, e0235226.	1.1	7
112	Human auditory ossicles as an alternative optimal source of ancient DNA. Genome Research, 2020, 30, 427-436.	2.4	37
113	Recovering signals of ghost archaic introgression in African populations. Science Advances, 2020, 6, eaax5097.	4.7	100
114	Identifying and Interpreting Apparent Neanderthal Ancestry in African Individuals. Cell, 2020, 180, 677-687.e16.	13.5	159
115	Ancient West African foragers in the context of African population history. Nature, 2020, 577, 665-670.	13.7	86
116	Genomic landscape of the signals of positive natural selection in populations of Northern Eurasia: A view from Northern Russia. PLoS ONE, 2020, 15, e0228778.	1.1	6
117	The genomic prehistory of peoples speaking Khoisan languages. Human Molecular Genetics, 2021, 30, R49-R55.	1.4	4
118	Origins of modern human ancestry. Nature, 2021, 590, 229-237.	13.7	166
119	The Main Nile Valley at the End of the Pleistocene (28–15Âka): Dispersal Corridor or Environmental Refugium?. Frontiers in Earth Science, 2021, 8, .	0.8	9
120	Continuity of the Middle Stone Age into the Holocene. Scientific Reports, 2021, 11, 70.	1.6	34
121	The deep population history in Africa. Human Molecular Genetics, 2021, 30, R2-R10.	1.4	15
122	Fine scale human genetic structure in three regions of Cameroon reveals episodic diversifying selection. Scientific Reports, 2021, 11, 1039.	1.6	3
123	Assessing the performance of qpAdm: a statistical tool for studying population admixture. Genetics, 2021, 217, .	1.2	84
124	A Pipeline for Assessment of Pathogenic Load in the Environment Using Microbiome Analysis. , 2021, , 493-510.		1
125	Structure and ancestry patterns of Ethiopians in genome-wide autosomal DNA. Human Molecular Genetics, 2021, 30, R42-R48.	1.4	3
126	Phenotypic diversity and history of the Congo Basin populations: Equatorial Guinea, Bantu Speaking Central Africans and African Pygmies. Annals of Human Biology, 2021, 48, 119-132.	0.4	0
127	Later Stone Age human hair from Vaalkrans Shelter, Cape Floristic Region of South Africa, reveals genetic affinity to Khoe groups. American Journal of Physical Anthropology, 2021, 174, 701-713.	2.1	3
128	Approaches to land snail shell bead manufacture in the Early Holocene of Malawi. Archaeological and Anthropological Sciences, 2021, 13, 1.	0.7	10

#	Article	IF	CITATIONS
129	Human burials at the Kisese <scp>II</scp> rockshelter, Tanzania. American Journal of Physical Anthropology, 2021, 175, 187-200.	2.1	5
132	Genetic diversity of the Sudanese: insights on origin and implications for health. Human Molecular Genetics, 2021, 30, R37-R41.	1.4	3
133	Human adaptation, demography and cattle domestication: an overview of the complexity of lactase persistence in Africa. Human Molecular Genetics, 2021, 30, R98-R109.	1.4	11
134	Small body size phenotypes among Middle and Later Stone Age Southern Africans. Journal of Human Evolution, 2021, 152, 102943.	1.3	6
135	The Middle to Later Stone Age transition at Panga ya Saidi, in the tropical coastal forest of eastern Africa. Journal of Human Evolution, 2021, 153, 102954.	1.3	18
136	Genetic substructure and complex demographic history of South African Bantu speakers. Nature Communications, 2021, 12, 2080.	5.8	47
137	Our Tangled Family Tree: New Genomic Methods Offer Insight into the Legacy of Archaic Admixture. Genome Biology and Evolution, 2021, 13, .	1.1	14
138	Earliest known human burial in Africa. Nature, 2021, 593, 95-100.	13.7	44
141	HSD17B7 gene in selfâ€renewal and oncogenicity of keratinocytes from Black versus White populations. EMBO Molecular Medicine, 2021, 13, e14133.	3.3	8
142	Iron Age hunting and herding in coastal eastern Africa: ZooMS identification of domesticates and wild bovids at Panga ya Saidi, Kenya. Journal of Archaeological Science, 2021, 130, 105368.	1.2	22
143	Human mobility at Tell Atchana (Alalakh), Hatay, Turkey during the 2nd millennium BC: Integration of isotopic and genomic evidence. PLoS ONE, 2021, 16, e0241883.	1.1	7
144	Evidence of the interplay of genetics and culture in Ethiopia. Nature Communications, 2021, 12, 3581.	5.8	25
146	Collagen fingerprinting traces the introduction of caprines to island Eastern Africa. Royal Society Open Science, 2021, 8, 202341.	1.1	10
147	Life in Deserts: The Genetic Basis of Mammalian Desert Adaptation. Trends in Ecology and Evolution, 2021, 36, 637-650.	4.2	35
148	<scp><i>Homo sapiens</i></scp> origins and evolution in the Kalahari Basin, southern Africa. Evolutionary Anthropology, 2021, 30, 327-344.	1.7	11
149	Insights into human history from the first decade of ancient human genomics. Science, 2021, 373, 1479-1484.	6.0	41
150	Peopling History of the Tibetan Plateau and Multiple Waves of Admixture of Tibetans Inferred From Both Ancient and Modern Genome-Wide Data. Frontiers in Genetics, 2021, 12, 725243.	1.1	27
152	African genetic diversity and adaptation inform a precision medicine agenda. Nature Reviews Genetics, 2021, 22, 284-306.	7.7	69

#	Article	IF	CITATIONS
153	Cultural extinction in evolutionary perspective. Evolutionary Human Sciences, 2021, 3, .	0.9	9
154	Ancient proteins provide evidence of dairy consumption in eastern Africa. Nature Communications, 2021, 12, 632.	5.8	39
155	Genomic History of Neolithic to Bronze Age Anatolia, Northern Levant, and Southern Caucasus. Cell, 2020, 181, 1158-1175.e28.	13.5	86
175	Asymmetric response of forest and grassy biomes to climate variability across the African Humid Period: influenced by anthropogenic disturbance?. Ecography, 2020, 43, 1118-1142.	2.1	16
176	Could There Have Been Human Families Where Parents Came from Different Populations: Denisovans, Neanderthals or Sapiens?. Scientia Et Fides, 2020, 8, 193.	0.3	1
177	An overview of the independent histories of the human Y chromosome and the human mitochondrial chromosome. The Proceedings of the International Conference on Creationism, 2018, 8, 133-151.	0.0	2
178	Recognizing plague epidemics in the archaeological record of West Africa. Afriques, 2018, , .	0.1	9
179	Revisiting the demographic history of Central African populations from a genetic perspective. , 0, , 1-29.		1
180	Genomes track Africa's historic populations. Nature, 2017, 549, 434-434.	13.7	0
181	East and Southern African Neolithic: Geography and Overview. , 2018, , 1-19.		0
189	The surgeon's role on chemical investigations of the composition of urinary stones. Urolithiasis, 2020, 48, 435-441.	1.2	1
191	East and Southern African Neolithic: Geography and Overview. , 2020, , 3537-3555.		Ο
192	Bantu-speaker migration and admixture in southern Africa. Human Molecular Genetics, 2021, 30, R56-R63.	1.4	21
197	Middle Stone Age Bifacial Technology and Pressure Flaking at the MIS 3 Site of Toumboura III, Eastern Senegal. African Archaeological Review, 2022, 39, 1-33.	0.8	9
198	Current Trends in Ancient DNA Study. , 2021, , 285-300.		0
200	Local ancestry prediction with <i>PyLAE</i> . PeerJ, 2021, 9, e12502.	0.9	0
201	Social stratification without genetic differentiation at the site of Kulubnarti in Christian Period Nubia. Nature Communications, 2021, 12, 7283.	5.8	13
202	Southern African foragers: Life histories of past communities. , 2022, , 145-159.		0

#	Article	IF	CITATIONS
203	Placing Ancient DNA Sequences into Reference Phylogenies. Molecular Biology and Evolution, 2022, 39, .	3.5	23
204	Southern African foragers: Geographical and historical context. , 2022, , 125-144.		0
205	Framework for osteobiographies: Eastern and Southern Africa. , 2022, , 99-110.		0
207	Genetics and Material Culture Support Repeated Expansions into Paleolithic Eurasia from a Population Hub Out of Africa. Genome Biology and Evolution, 2022, 14, .	1.1	15
208	Ancient DNA and deep population structure in sub-Saharan African foragers. Nature, 2022, 603, 290-296.	13.7	51
209	A unified genealogy of modern and ancient genomes. Science, 2022, 375, eabi8264.	6.0	59
210	Ancient DNA illuminates how humans travelled and interacted in Stone Age Africa. Nature, 2022, , .	13.7	0
211	Hunter-gatherer genomes reveal diverse demographic trajectories during the rise of farming in Eastern Africa. Current Biology, 2022, 32, 1852-1860.e5.	1.8	15
213	South-to-north migration preceded the advent of intensive farming in the Maya region. Nature Communications, 2022, 13, 1530.	5.8	21
214	Ancient genomes from the Himalayas illuminate the genetic history of Tibetans and their Tibeto-Burman speaking neighbors. Nature Communications, 2022, 13, 1203.	5.8	25
215	Coring, profiling, and trenching: Archaeological field strategies for investigating the Pleistocene-Holocene-Anthropocene continuum. Quaternary International, 2022, 628, 1-17.	0.7	7
216	Hunter-gatherer technological organization and responses to Holocene climate change in coastal, lakeshore, and grassland ecologies of eastern Africa. Quaternary Science Reviews, 2022, 280, 107390.	1.4	7
218	Male-biased migration from East Africa introduced pastoralism into southern Africa. BMC Biology, 2021, 19, 259.	1.7	4
219	Ostrich eggshell beads reveal 50,000-year-old social network in Africa. Nature, 2022, 601, 234-239.	13.7	32
221	Apportioning archaic variants among modern populations. Philosophical Transactions of the Royal Society B: Biological Sciences, 2022, 377, 20200411.	1.8	11
222	Exome sequencing of families from Ghana reveals known and candidate hearing impairment genes. Communications Biology, 2022, 5, 369.	2.0	8
236	Human evolution: The unsealed fates of foragers andÂfarmers. Current Biology, 2022, 32, R362-R365.	1.8	0
240	Genomic Insights Into the Demographic History of the Southern Chinese. Frontiers in Ecology and Evolution, 0, 10, .	1.1	13

#	ARTICLE Phylogeography of Sub-Saharan Mitochondrial Lineages Outside Africa Highlights the Roles of the Holocope Climate Changes and the Atlantic Slave Trade. International Journal of Molecular Sciences	IF	CITATIONS
243	2022, 23, 9219.	1.8	2
244	European Trade in Malawi: The Glass Bead Evidence. African Archaeological Review, 2023, 40, 377-396.	0.8	1
246	Hominin Evolution II. , 2022, , 50-63.		0
247	Direct detection of natural selection in Bronze Age Britain. Genome Research, 2022, 32, 2057-2067.	2.4	20
248	The petrous bone contains high concentrations of osteocytes: One possible reason why ancient DNA is better preserved in this bone. PLoS ONE, 2022, 17, e0269348.	1.1	6
249	Blue Turns to Gray: Paleogenomic Insights into the Evolutionary History and Extinction of the Blue Antelope ( <i>Hippotragus leucophaeus</i> ). Molecular Biology and Evolution, 2022, 39, .	3.5	5
250	Pleistocene parades of carnivores into North America. Molecular Ecology, 0, , .	2.0	1
251	4000-year-old hair from the Middle Nile highlights unusual ancient DNA degradation pattern and a potential source of early eastern Africa pastoralists. Scientific Reports, 2022, 12, .	1.6	1
252	The Hofmeyr Bony Labyrinth: Morphological Description and Affinity. Vertebrate Paleobiology and Paleoanthropology, 2022, , 165-178.	0.1	1
253	Genetic Divergence Within Southern Africa During the Later Stone Age. Vertebrate Paleobiology and Paleoanthropology, 2022, , 19-28.	0.1	2
254	The Dentition of the Hofmeyr Skull. Vertebrate Paleobiology and Paleoanthropology, 2022, , 213-233.	0.1	1
255	Genomic perspectives on human dispersals during the Holocene. Proceedings of the National Academy of Sciences of the United States of America, 2023, 120, .	3.3	12
257	The first out of Africa migrants. , 2023, , 361-395.		0
259	Whole-genome sequencing reveals a complex African population demographic history and signatures of local adaptation. Cell, 2023, 186, 923-939.e14.	13.5	34
260	Small game forgotten: Late Pleistocene foraging strategies in eastern Africa, and remote capture at Panga ya Saidi, Kenya. Quaternary Science Reviews, 2023, 305, 108032.	1.4	3
261	Human genetic history on the Tibetan Plateau in the past 5100 years. Science Advances, 2023, 9, .	4.7	17
262	Evolutionary Genetics and Admixture in African Populations. Genome Biology and Evolution, 2023, 15, .	1.1	7
263	Entwined African and Asian genetic roots of medieval peoples of the Swahili coast. Nature, 2023, 615, 866-873.	13.7	17

#	Article	IF	CITATIONS
265	Dense sampling of ethnic groups within African countries reveals fine-scale genetic structure and extensive historical admixture. Science Advances, 2023, 9, .	4.7	7
274	Africa, Tropical: East African Foragers. , 2024, , 126-131.		0
280	Panga ya Saidi, Kenya. , 2023, , 665-672.		0