Genetic evidence for archaic admixture in Africa

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Citation Report

#	Article	IF	Citations
1	Next-generation sequencing technologies and applications for human genetic history and forensics. Investigative Genetics, 2011, 2, 23.	3.3	101
2	Paleogenomics of Archaic Hominins. Current Biology, 2011, 21, R1002-R1009.	1.8	39
3	Modeling Human Ecodynamics and Biocultural Interactions in the Late Pleistocene of Western Eurasia. Human Ecology, 2011, 39, 705-725.	0.7	82
4	A New Isolation with Migration Model along Complete Genomes Infers Very Different Divergence Processes among Closely Related Great Ape Species. PLoS Genetics, 2012, 8, e1003125.	1.5	102
5	Genomic Data Reveal a Complex Making of Humans. PLoS Genetics, 2012, 8, e1002837.	1.5	43
6	Archaic human genomics. American Journal of Physical Anthropology, 2012, 149, 24-39.	2.1	19
7	Genomic Variation in Seven Khoe-San Groups Reveals Adaptation and Complex African History. Science, 2012, 338, 374-379.	6.0	364
8	Human Evolution Out of Africa: The Role of Refugia and Climate Change. Science, 2012, 335, 1317-1321.	6.0	239
9	Evolutionary History and Adaptation from High-Coverage Whole-Genome Sequences of Diverse African Hunter-Gatherers. Cell, 2012, 150, 457-469.	13.5	289
10	Longer time scale for human evolution. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 15531-15532.	3.3	15
11	Effect of ancient population structure on the degree of polymorphism shared between modern human populations and ancient hominins. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 13956-13960.	3.3	207
12	Paleopopulation Genetics. Annual Review of Genetics, 2012, 46, 635-649.	3.2	17
13	Accurate whole-genome sequencing and haplotyping from 10 to 20 human cells. Nature, 2012, 487, 190-195.	13.7	226
14	Revising the human mutation rate: implications for understanding human evolution. Nature Reviews Genetics, 2012, 13, 745-753.	7.7	483
15	A Haplotype at STAT2 Introgressed from Neanderthals and Serves as a Candidate of Positive Selection in Papua New Guinea. American Journal of Human Genetics, 2012, 91, 265-274.	2.6	152
16	Inland human settlement in southern Arabia 55,000 years ago. New evidence from the Wadi Surdud Middle Paleolithic site complex, western Yemen. Journal of Human Evolution, 2012, 63, 452-474.	1.3	102
18	Population structure and migration in Africa: correlations between archaeological, linguistic, and genetic data. , 0, , 135-171.		2
19	Why do we migrate? A retrospective. , 2012, , 527-536.		4

#	Article	IF	Citations
20	Hunter-gatherer genomes a trove of genetic diversity. Nature, 2012, , .	13.7	0
21	Science and Medicine. Yearbook of Paediatric Endocrinology, 2012, , 223-237.	0.0	0
22	Sympatric Speciation in the Post â€œModern Synthesisâ€•Era of Evolutionary Biology. Evolutionary Biology, 2012, 39, 158-180.	0.5	89
23	The environmental context for the origins of modern human diversity: A synthesis of regional variability in African climate 150,000â€“30,000 years ago. Journal of Human Evolution, 2012, 62, 563-592.	1.3	240
24	What makes a modern human. Nature, 2012, 485, 33-35.	13.7	61
25	The status of Homo heidelbergensis (Schoetensack 1908). Evolutionary Anthropology, 2012, 21, 101-107.	1.7	270
26	Nothing in medicine makes sense, except in the light of evolution. Journal of Molecular Medicine, 2012, 90, 481-494.	1.7	38
27	Significance of Neandertal and Denisovan Genomes in Human Evolution. Annual Review of Anthropology, 2013, 42, 433-449.	0.4	34
28	Selection and Adaptation in the Human Genome. Annual Review of Genomics and Human Genetics, 2013, 14, 467-489.	2.5	116
33	Possible origins and evolution of the hepatitis B virus (HBV). Seminars in Cancer Biology, 2013, 23, 561-575.	4.3	76
34	Population Genomics of Human Adaptation. Annual Review of Ecology, Evolution, and Systematics, 2013, 44, 123-143.	3.8	81
35	Phylogenetic Relationships (Biomolecules). , 2013, , 1-25.		0
36	Higher Levels of Neanderthal Ancestry in East Asians than in Europeans. Genetics, 2013, 194, 199-209.	1.2	219
38	Nubian Complex reduction strategies in Dhofar, southern Oman. Quaternary International, 2013, 300, 244-266.	0.7	73
39	Agreements and Misunderstandings among Three Scientific Fields. Current Anthropology, 2013, 54, S214-S220.	0.8	10
40	Variability in the Middle Stone Age of Eastern Africa. Current Anthropology, 2013, 54, S234-S254.	0.8	151
41	An African American Paternal Lineage Adds an Extremely Ancient Root to the Human Y Chromosome Phylogenetic Tree. American Journal of Human Genetics, 2013, 92, 454-459.	2.6	124
42	Hominin evolution and gene flow in the Pleistocene Africa. Anthropologischer Anzeiger, 2013, 70, 221-227.	0.2	3

#	Article	IF	Citations
43	Robust Demographic Inference from Genomic and SNP Data. PLoS Genetics, 2013, 9, e1003905.	1.5	1,185
44	Bridging disciplines to better elucidate the evolution of early Homo sapiens in southern Africa. South African Journal of Science, 2013, 109, 8.	0.3	15
45	Geologic Life: Prehistory, Climate, Futures in the Anthropocene. Environment and Planning D: Society and Space, 2013, 31, 779-795.	2.3	257
46	Apparent Variation in Neanderthal Admixture among African Populations is Consistent with Gene Flow from Non-African Populations. Genome Biology and Evolution, 2013, 5, 2075-2081.	1.1	31
47	Nuclear Genetic Diversity in Human Lice (Pediculus humanus) Reveals Continental Differences and High Inbreeding among Worldwide Populations. PLoS ONE, 2013, 8, e57619.	1.1	46
48	X-Linked MTMR8 Diversity and Evolutionary History of Sub-Saharan Populations. PLoS ONE, 2013, 8, e80710.	1.1	1
49	To name or not to name: Criteria to promote economy of change in Linnaean classification schemes. Zootaxa, 2013, 3636, 201-44.	0.2	170
50	Archaeogenetics of Africa and of the African Hunter-Gatherers. , 2014, , .		0
51	Human evolution. , 0, , 13-28.		0
52	Why we are not all multiregionalists now. Trends in Ecology and Evolution, 2014, 29, 248-251.	4.2	57
53	The place of the Neanderthals in hominin phylogeny. Journal of Anthropological Archaeology, 2014, 35, 32-50.	0.7	12
54	Resurrecting Surviving Neandertal Lineages from Modern Human Genomes. Science, 2014, 343, 1017-1021.	6.0	510
55	The impact of whole-genome sequencing on the reconstruction of human population history. Nature Reviews Genetics, 2014, 15, 149-162.	7.7	147
56	Human Evolution: Genomic Gifts from Archaic Hominins. Current Biology, 2014, 24, R845-R848.	1.8	8
57	DiagnosingHomo sapiensin the fossil record. Annals of Human Biology, 2014, 41, 312-322.	0.4	28
58	The peopling of the African continent and the diaspora into the new world. Current Opinion in Genetics and Development, 2014, 29, 120-132.	1.5	45
59	Population-specific common SNPs reflect demographic histories and highlight regions of genomic plasticity with functional relevance. BMC Genomics, 2014, 15, 437.	1.2	40
60	Genetic Variation and Adaptation in Africa: Implications for Human Evolution and Disease. Cold Spring Harbor Perspectives in Biology, 2014, 6, a008524-a008524.	2.3	87

#	Article	IF	Citations
61	Iwo Eleru's place among Late Pleistocene and Early Holocene populations of North and East Africa. Journal of Human Evolution, 2014, 75, 80-89.	1.3	22
62	Molecular Editing of Cellular Responses by the High-Affinity Receptor for IgE. Science, 2014, 343, 1021-1025.	6.0	81
63	Early humans: tools, language, and culture. , 2015, , 339-361.		1
64	Early modern human dispersal from Africa: genomic evidence for multiple waves of migration. Investigative Genetics, 2015, 6, 13.	3.3	34
65	Next-generation sequencing and the expanding domain of phylogeography. Folia Zoologica, 2015, 64, 187-206.	0.9	47
66	Africa from 48,000 to 9500BCE. , 2015, , 362-393.		21
67	A Hominin Femur with Archaic Affinities from the Late Pleistocene of Southwest China. PLoS ONE, 2015, 10, e0143332.	1.1	13
68	Neanderthals had outsize effect on human biology. Nature, 2015, 523, 512-513.	13.7	4
69	The Evolution and Functional Impact of Human Deletion Variants Shared with Archaic Hominin Genomes. Molecular Biology and Evolution, 2015, 32, 1008-1019.	3.5	45
70	Interleukin-37 gene variants segregated anciently coexist during hominid evolution. European Journal of Human Genetics, 2015, 23, 1392-1398.	1.4	14
71	Reticulate Evolution. Interdisciplinary Evolution Research, 2015, , .	0.2	19
72	Testing modern human out-of-Africa dispersal models and implications for modern human origins. Journal of Human Evolution, 2015, 87, 95-106.	1.3	58
73	Worldwide Population Structure, Long-Term Demography, and Local Adaptation of Helicobacter pylori. Genetics, 2015, 200, 947-963.	1.2	65
74	Divergence-with-Gene-Flowâ€"What Humans and Other Mammals Got up to. Interdisciplinary Evolution Research, 2015, , 255-295.	0.2	2
75	When mothers need others: The impact of hominin life history evolution on cooperative breeding. Journal of Human Evolution, 2015, 84, 16-24.	1.3	38
76	Evidence for archaic adaptive introgression in humans. Nature Reviews Genetics, 2015, 16, 359-371.	7.7	471
77	Bias in estimators of archaic admixture. Theoretical Population Biology, 2015, 100, 63-78.	0.5	25
78	An Evolutionary Anthropological Perspective on Modern Human Origins. Annual Review of Anthropology, 2015, 44, 533-556.	0.4	83

#	Article	IF	Citations
79	Small Amounts of Archaic Admixture Provide Big Insights into Human History. Cell, 2015, 163, 281-284.	13.5	53
80	Ancient Ethiopian genome reveals extensive Eurasian admixture in Eastern Africa. Science, 2015, 350, 820-822.	6.0	277
81	IBD Sharing between Africans, Neandertals, and Denisovans. Genome Biology and Evolution, 2016, 8, 3406-3416.	1.1	6
83	CoMuS: simulating coalescent histories and polymorphic data from multiple species. Molecular Ecology Resources, 2016, 16, 1435-1448.	2.2	9
84	Sexual Deviance and Society. , 0, , .		11
85	The Importance of Croatian Pleistocene Hominin Finds in the Study of Human Evolution. Vertebrate Paleobiology and Paleoanthropology, 2016, , 35-50.	0.1	7
86	Late Stone Age human remains from Ishango (Democratic RepublicÂofÂCongo): New insights on Late Pleistocene modernÂhumanÂdiversity in Africa. Journal of Human Evolution, 2016, 96, 35-57.	1.3	34
87	Archaic admixture in human history. Current Opinion in Genetics and Development, 2016, 41, 93-97.	1.5	26
89	Genomic analysis of Andamanese provides insights into ancient human migration into Asia and adaptation. Nature Genetics, 2016, 48, 1066-1070.	9.4	126
90	Inferences of African evolutionary history from genomic data. Current Opinion in Genetics and Development, 2016, 41, 159-166.	1.5	34
91	Ancient DNA and human history. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 6380-6387.	3.3	178
92	The Middle Stone Age archaeology of the Senegal River Valley. Quaternary International, 2016, 408, 16-32.	0.7	27
93	A demographic perspective on the Middle to Later Stone Age transition from Nasera rockshelter, Tanzania. Philosophical Transactions of the Royal Society B: Biological Sciences, 2016, 371, 20150238.	1.8	79
94	Detecting hybridization using ancient <scp>DNA</scp>. Molecular Ecology, 2016, 25, 2398-2412.	2.0	37
95	Origins and Evolution of Hepatitis B Virus and Hepatitis D Virus. Cold Spring Harbor Perspectives in Medicine, 2016, 6, a021360.	2.9	84
96	Whole-genome sequence analyses of Western Central African Pygmy hunter-gatherers reveal a complex demographic history and identify candidate genes under positive natural selection. Genome Research, 2016, 26, 279-290.	2.4	54
97	Model-based analyses of whole-genome data reveal a complex evolutionary history involving archaic introgression in Central African Pygmies. Genome Research, 2016, 26, 291-300.	2.4	87
99	Africa from MIS 6-2: The Florescence of Modern Humans. Vertebrate Paleobiology and Paleoanthropology, 2016, , 1-20.	0.1	8

#	Article	IF	Citations
100	The Later Pleistocene in the Northeastern Central African Rainforest. Vertebrate Paleobiology and Paleoanthropology, 2016, , 301-319.	0.1	10
101	The Late Quaternary Hominins of Africa: The Skeletal Evidence from MIS 6-2. Vertebrate Paleobiology and Paleoanthropology, 2016, , 323-381.	0.1	31
102	The Middle Palaeolithic of West Africa: Lithic techno-typological analyses of the site of Tiemassas, Senegal. Quaternary International, 2016, 408, 4-15.	0.7	15
103	The Hybrid Origin of â€œModernâ€•Humans. Evolutionary Biology, 2016, 43, 1-11.	0.5	117
104	Tracing the peopling of the world through genomics. Nature, 2017, 541, 302-310.	13.7	562
105	Persistence of Middle Stone Age technology to the Pleistocene/Holocene transition supports a complex hominin evolutionary scenario in West Africa. Journal of Archaeological Science: Reports, 2017, 11, 639-646.	0.2	15
106	Manot 1 calvaria and Recent Modern Human Evolution: an Anthropological Perspective. Bulletins Et Memoires De La Societe D'Anthropologie De Paris, 2017, 29, 119-130.	0.0	12
107	Migrating microbes: what pathogens can tell us about population movements and human evolution. Annals of Human Biology, 2017, 44, 397-407.	0.4	22
108	Modelling the role of groundwater hydro-refugia in East African hominin evolution and dispersal. Nature Communications, 2017, 8, 15696.	5.8	47
109	Ancient oncogenesis, infection and human evolution. Evolutionary Applications, 2017, 10, 949-964.	1.5	15
110	Living in an adaptive world: Genomic dissection of the genus Homo and its immune response. Journal of Experimental Medicine, 2017, 214, 877-894.	4.2	34
111	Southern African ancient genomes estimate modern human divergence to 350,000 to 260,000 years ago. Science, 2017, 358, 652-655.	6.0	351
112	Reconstructing Prehistoric African Population Structure. Cell, 2017, 171, 59-71.e21.	13.5	308
113	Variation and Functional Impact of Neanderthal Ancestry in Western Asia. Genome Biology and Evolution, 2017, 9, 3516-3524.	1.1	14
114	The contribution of admixture to primate evolution. Current Opinion in Genetics and Development, 2017, 47, 61-68.	1.5	44
115	Archaic Hominin Introgression in Africa Contributes to Functional Salivary MUC7 Genetic Variation. Molecular Biology and Evolution, 2017, 34, 2704-2715.	3.5	57
117	The Contribution of Genetic Ancestry From Archaic Humans to Modern Humans. , 2017, , 55-63.		0
118	Insights into Modern Human Prehistory Using Ancient Genomes. Trends in Genetics, 2018, 34, 184-196.	2.9	50

#	Article	IF	Citations
119	Craniomandibular form and body size variation of first generation mouse hybrids: A model for hominin hybridization. Journal of Human Evolution, 2018, 116, 57-74.	1.3	15
120	Tales of Human Migration, Admixture, and Selection in Africa. Annual Review of Genomics and Human Genetics, 2018, 19, 405-428.	2.5	78
121	Analysis of Human Sequence Data Reveals Two Pulses of Archaic Denisovan Admixture. Cell, 2018, 173, 53-61.e9.	13.5	271
122	Between continuity and discontinuity: An overview of the West African Paleolithic over the last 200,000 years. Quaternary International, 2018, 466, 3-22.	0.7	22
123	Clarifying distinct models of modern human origins in Africa. Current Opinion in Genetics and Development, 2018, 53, 148-156.	1.5	51
124	Outstanding questions in the study of archaic hominin admixture. PLoS Genetics, 2018, 14, e1007349.	1.5	50
125	Cultural Exaptation and Cultural Neural Reuse: A Mechanism for the Emergence of Modern Culture and Behavior. Biological Theory, 2018, 13, 213-227.	0.8	30
126	Evolutionary and Medical Consequences of Archaic Introgression into Modern Human Genomes. Genes, 2018, 9, 358.	1.0	28
127	â€˜Behavioral modernityâ€™ as a process, not an event, in the human niche. Time and Mind, 2018, 11, 163-183.	0.4	21
128	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
129	Introduction: Interrogating the Anthropocene. , 2018, , 1-71.		4
130	Phylogeny Estimation by Integration over Isolation with Migration Models. Molecular Biology and Evolution, 2018, 35, 2805-2818.	3.5	89
131	The demographic and adaptive history of central African hunter-gatherers and farmers. Current Opinion in Genetics and Development, 2018, 53, 90-97.	1.5	17
132	Early Hominins. , 2018, , 33-60.		0
133	The Settlement of the Near East. , 2018, , 133-174.		0
135	A statistical model for reference-free inference of archaic local ancestry. PLoS Genetics, 2019, 15, e1008175.	1.5	31
136	No Country for Oldowan Men: Emerging Factors in Language Evolution. Frontiers in Psychology, 2019, 10, 1448.	1.1	12
137	Philosophical Urbanism. , 2019, , .		2

#	Article	IF	Citations
138	A method for genome-wide genealogy estimation for thousands of samples. Nature Genetics, 2019, 51, 1321-1329.	9.4	338
139	Genomics of disease risk in globally diverse populations. Nature Reviews Genetics, 2019, 20, 520-535.	7.7	217
140	The Emergence of Humanity. , 2019, , 399-470.		0
141	Species-specific effects of climate change on the distribution of suitable baboon habitats â€“ Ecological niche modeling of current and Last Glacial Maximum conditions. Journal of Human Evolution, 2019, 132, 215-226.	1.3	28
142	Searching for archaic contribution in Africa. Annals of Human Biology, 2019, 46, 129-139.	0.4	4
143	Models of archaic admixture and recent history from two-locus statistics. PLoS Genetics, 2019, 15, e1008204.	1.5	57
144	Ancient admixture from an extinct ape lineage into bonobos. Nature Ecology and Evolution, 2019, 3, 957-965.	3.4	59
145	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
146	Multiple Deeply Divergent Denisovan Ancestries in Papuans. Cell, 2019, 177, 1010-1021.e32.	13.5	181
147	Identification of African-Specific Admixture between Modern and Archaic Humans. American Journal of Human Genetics, 2019, 105, 1254-1261.	2.6	16
148	Archaic hominin introgression into modern human genomes. American Journal of Physical Anthropology, 2020, 171, 60-73.	2.1	33
149	Neutral evolution of human enamelâ€“dentine junction morphology. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 26183-26189.	3.3	11
150	The Middle Stone Age occupations of TiÃ©massas, coastal West Africa, between 62 and 25 thousand years ago. Journal of Archaeological Science: Reports, 2020, 34, 102658.	0.2	3
151	Mapping gene flow between ancient hominins through demography-aware inference of the ancestral recombination graph. PLoS Genetics, 2020, 16, e1008895.	1.5	76
152	On the Operation of Retouch in Southern Africaâ€™s Early Middle Stone Age. Journal of Paleolithic Archaeology, 2020, 3, 1149-1179.	0.7	2
153	The reversal of human phylogeny: Homo left Africa as erectus, came back as sapiens sapiens. Hereditas, 2020, 157, 51.	0.5	3
154	VolcanoFinder: Genomic scans for adaptive introgression. PLoS Genetics, 2020, 16, e1008867.	1.5	62
155	Toward capturing the functional and nuanced nature of social stereotypes: An affordance management approach. Advances in Experimental Social Psychology, 2020, 62, 245-304.	2.0	21

#	Article	IF	Citations
156	A massacre of early Neolithic farmers in the high Pyrenees at Els Trocs, Spain. Scientific Reports, 2020, 10, 2131.	1.6	20
157	Recovering signals of ghost archaic introgression in African populations. Science Advances, 2020, 6, eaax5097.	4.7	100
158	Ancient West African foragers in the context of African population history. Nature, 2020, 577, 665-670.	13.7	86
159	HBV evolution and genetic variability: Impact on prevention, treatment and development of antivirals. Antiviral Research, 2021, 186, 104973.	1.9	28
160	The genomic prehistory of peoples speaking Khoisan languages. Human Molecular Genetics, 2021, 30, R49-R55.	1.4	4
161	Origins of modern human ancestry. Nature, 2021, 590, 229-237.	13.7	166
162	Population Variation of the Human Genome. , 2021, , 329-350.		0
163	Constraining the Likely Technological Niches of Late Middle Pleistocene Hominins with Homo naledi as Case Study. Journal of Archaeological Method and Theory, 2021, 28, 11-52.	1.4	8
164	Population Genomics of High-Altitude Adaptation. Evolutionary Studies, 2021, , 67-100.	0.2	0
165	The influence of evolutionary history on human health and disease. Nature Reviews Genetics, 2021, 22, 269-283.	7.7	133
166	The deep population history in Africa. Human Molecular Genetics, 2021, 30, R2-R10.	1.4	15
167	Inferring Human Demographic History from Genetic Data. , 2021, , 187-204.		0
168	Fine scale human genetic structure in three regions of Cameroon reveals episodic diversifying selection. Scientific Reports, 2021, 11, 1039.	1.6	3
169	Our Tangled Family Tree: New Genomic Methods Offer Insight into the Legacy of Archaic Admixture. Genome Biology and Evolution, 2021, 13, .	1.1	14
170	Authorship Patterns in Cancer Genomics Publications Across Africa. JCO Global Oncology, 2021, 7, 747-755.	0.8	8
171	Inferring archaic introgression from hominin genetic data. Evolutionary Anthropology, 2021, 30, 199-220.	1.7	9
173	Origins and Evolution of the Primate Hepatitis B Virus. Frontiers in Microbiology, 2021, 12, 653684.	1.5	14
174	The history and evolution of the Denisovan- EPAS1 haplotype in Tibetans. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	3.3	48

#	Article	IF	Citations
175	An ancestral recombination graph of human, Neanderthal, and Denisovan genomes. Science Advances, 2021, 7, .	4.7	47
176	<scp>Homo sapiens</scp>origins and evolution in the Kalahari Basin, southern Africa. Evolutionary Anthropology, 2021, 30, 327-344.	1.7	11
177	Philippine Ayta possess the highest level of Denisovan ancestry in the world. Current Biology, 2021, 31, 4219-4230.e10.	1.8	37
179	The ripples of modernity: How we can extend paleoanthropology with the extended evolutionary synthesis. Evolutionary Anthropology, 2021, 30, 84-98.	1.7	14
181	Neanderthals and Their Contemporaries. , 2015, , 2243-2279.		9
182	Human ancestors interbred with related species. Nature, 0, , .	13.7	1
192	Introgression Makes Waves in Inferred Histories of Effective Population Size. Human Biology, 2017, 89, 67.	0.4	14
193	The Later Stone Age Calvaria from Iwo Eleru, Nigeria: Morphology and Chronology. PLoS ONE, 2011, 6, e24024.	1.1	107
194	The Nubian Complex of Dhofar, Oman: An African Middle Stone Age Industry in Southern Arabia. PLoS ONE, 2011, 6, e28239.	1.1	172
195	North African Populations Carry the Signature of Admixture with Neandertals. PLoS ONE, 2012, 7, e47765.	1.1	67
196	Neanderthal and Denisova tooth protein variants in present-day humans. PLoS ONE, 2017, 12, e0183802.	1.1	15
197	Spatial and Temporal Simulation of Human Evolution. Methods, Frameworks and Applications. Current Genomics, 2014, 15, 245-255.	0.7	16
198	Reconsideration of the â€œOut of Africaâ€•Concept as Not Having Enough Proof. Advances in Anthropology, 2014, 04, 18-37.	0.1	9
199	Homo naledi and Pleistocene hominin evolution in subequatorial Africa. ELife, 2017, 6, .	2.8	75
200	Haplotypes spanning centromeric regions reveal persistence of large blocks of archaic DNA. ELife, 2019, 8, .	2.8	54
201	Embracing heterogeneity: coalescing the Tree of Life and the future of phylogenomics. PeerJ, 2019, 7, e6399.	0.9	111
202	Revisiting the demographic history of Central African populations from a genetic perspective. , 0, , 1-29.		1
205	Neanderthals and Their Contemporaries. , 2014, , 1-35.		2

#	Article	IF	Citations
206	Phylogenetic Relationships of Hominids: Biomolecular Approach. , 2015, , 2015-2041.		0
211	UneÂperspective gÃ©nÃ©tique surÂnotre histoireÂ: migrations humaines etÂadaptation ÃÂlâ€™environnement. , 2017, , 33-60.		0
212	Identical by Descent (IBD): Investigation of the Genetic Ties between Africans, Denisovans, and Neandertals. Asian Journal of Humanity Art and Literature, 2017, 4, 157-170.	0.2	8
213	Transition to Agriculture and First State Presence: A Global Analysis. SSRN Electronic Journal, 0, , .	0.4	2
214	Sky and Gender Myths in the Founding of Early Built Environments. , 2019, , 43-70.		0
216	Variation in Middle Stone Age mandibular molar enamel-dentine junction topography at Klasies River Main Site assessed by diffeomorphic surface matching. Journal of Human Evolution, 2021, 161, 103079.	1.3	4
218	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
220	High Genetic Diversity and Rickettsia felis in Pediculus humanus Lice Infesting Mbuti (pygmy people), -Democratic Republic of Congo. Frontiers in Cellular and Infection Microbiology, 2022, 12, 834388.	1.8	3
221	The Syngameon Enigma. Plants, 2022, 11, 895.	1.6	11
222	Tracing of Human Migration and Diversity by Forensic DNA Analysis. , 2022, , 1165-1184.		0
223	Paleogenomics, hominin interbreeding and language evolution. Journal of Anthropological Sciences, 2013, 91, 239-44.	0.4	2
224	Archaic human genomes and language evolution. Journal of Anthropological Sciences, 2013, 91, 253-5.	0.4	0
225	A review of the spread and habitat of the genus <i>Homo</i>:. Anthropological Science, 2022, , .	0.2	0
226	Mitochondrial Pseudogenes Suggest Repeated Inter-Species Hybridization among Direct Human Ancestors. Genes, 2022, 13, 810.	1.0	5
232	Panâ€Africanism vs. singleâ€origin of Homo sapiens: Putting the debate in the light of evolutionary biology. Evolutionary Anthropology, 2022, 31, 199-212.	1.7	6
233	Midfacial Morphology and Neandertalâ€“Modern Human Interbreeding. Biology, 2022, 11, 1163.	1.3	2
234	No Homo: Why Theistic Evolution Fails. , 2022, 2, 26-34.		0
235	Merging morphological and genetic evidence to assess hybridization in Western Eurasian late Pleistocene hominins. Nature Ecology and Evolution, 2022, 6, 1573-1585.	3.4	12

#	Article	IF	Citations
236	A Pleistocene assemblage of nearâ€modern Papio hamadryas from the Middle Awash study area, Afar Rift, Ethiopia. American Journal of Biological Anthropology, 0, , .	0.6	3
237	Moving beyond the adaptationist paradigm for human evolution, and why it matters. Journal of Human Evolution, 2023, 174, 103296.	1.3	5
238	The Emergence of Habitual Ochre Use in Africa and its Significance for The Development of Ritual Behavior During The Middle Stone Age. Journal of World Prehistory, 2022, 35, 233-319.	1.1	14
239	Is there still evolution in the human population?. Biologia Futura, 2022, 73, 359-374.	0.6	1
240	Hominin fossils: Anatomical trends. , 2023, , 165-217.		0
241	A Population Genetic Perspective on Subsistence Systems in the Sahel/Savannah Belt of Africa and the Historical Role of Pastoralism. Genes, 2023, 14, 758.	1.0	1
242	The Position of the Hofmeyr Skull within Late Pleistocene and Holocene African Regional Diversity: 2D and 3D Morphometric Analyses. Vertebrate Paleobiology and Paleoanthropology, 2022, , 119-141.	0.1	5
243	Introduction: The Fossil Record of Homo sapiens in Africa â€“ Morphological Variability in the Late Quaternary and the Significance of the Hofmeyr Skull. Vertebrate Paleobiology and Paleoanthropology, 2022, , 1-5.	0.1	0
244	Modern Humans Disperse From Africa. , 2022, , 581-623.		0
248	An African-wide origin of Homo sapiens. , 2023, , 331-360.		0
249	Interwoven evolution of Homo erectus, Homo floresiensis, and Homo sapiens. , 2023, , 551-573.		0
251	Biosocial complexity and the skull. , 2023, , 39-72.		0
252	Evolution in biomechanics. , 2023, , 495-663.		0
253	Evolutionary Genetics and Admixture in African Populations. Genome Biology and Evolution, 2023, 15, .	1.1	7
254	On the limits of fitting complex models of population history to f-statistics. ELife, 0, 12, .	2.8	35
259	IhÃ² ElÃ©Ã©rÃº [Iwo Eleru], Nigeria. , 2023, , 915-925.		0