

Mark G Thomas

List of Publications by Year in descending order

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Version: 2024-02-01

180
papers

20,621
citations

10986
71
h-index

11607
135
g-index

193
all docs

193
docs citations

193
times ranked

18092
citing authors

#	ARTICLE	IF	CITATIONS
1	Ancient human genomes suggest three ancestral populations for present-day Europeans. <i>Nature</i> , 2014, 513, 409-413.	27.8	1,179
2	Late Pleistocene Demography and the Appearance of Modern Human Behavior. <i>Science</i> , 2009, 324, 1298-1301.	12.6	952
3	Tracing European Founder Lineages in the Near Eastern mtDNA Pool. <i>American Journal of Human Genetics</i> , 2000, 67, 1251-1276.	6.2	837
4	Current perspectives and the future of domestication studies. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 6139-6146.	7.1	594
5	Regional population collapse followed initial agriculture booms in mid-Holocene Europe. <i>Nature Communications</i> , 2013, 4, 2486.	12.8	532
6	The Beaker phenomenon and the genomic transformation of northwest Europe. <i>Nature</i> , 2018, 555, 190-196.	27.8	503
7	A genomic history of Aboriginal Australia. <i>Nature</i> , 2016, 538, 207-214.	27.8	439
8	Genetic Discontinuity Between Local Hunter-Gatherers and Central Europe's First Farmers. <i>Science</i> , 2009, 326, 137-140.	12.6	433
9	Population genetic structure of variable drug response. <i>Nature Genetics</i> , 2001, 29, 265-269.	21.4	425
10	Absence of the lactase-persistence-associated allele in early Neolithic Europeans. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 3736-3741.	7.1	406
11	The Origins of Lactase Persistence in Europe. <i>PLoS Computational Biology</i> , 2009, 5, e1000491.	3.2	383
12	Early farmers from across Europe directly descended from Neolithic Aegeans. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 6886-6891.	7.1	376
13	Lactose digestion and the evolutionary genetics of lactase persistence. <i>Human Genetics</i> , 2009, 124, 579-591.	3.8	367
14	Genomic analyses inform on migration events during the peopling of Eurasia. <i>Nature</i> , 2016, 538, 238-242.	27.8	360
15	A recent bottleneck of Y chromosome diversity coincides with a global change in culture. <i>Genome Research</i> , 2015, 25, 459-466.	5.5	348
16	Evolution of lactase persistence: an example of human niche construction. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2011, 366, 863-877.	4.0	340
17	Did Our Species Evolve in Subdivided Populations across Africa, and Why Does It Matter?. <i>Trends in Ecology and Evolution</i> , 2018, 33, 582-594.	8.7	315
18	Reconstructing Prehistoric African Population Structure. <i>Cell</i> , 2017, 171, 59-71.e21.	28.9	308

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19	A new time-scale for ray-finned fish evolution. Proceedings of the Royal Society B: Biological Sciences, 2007, 274, 489-498.	2.6	298
20	Presence of a newly described human DNA virus (TTV) in patients with liver disease. Lancet, The, 1998, 352, 195-197.	13.7	297
21	Rethinking the dispersal of <i>Homo sapiens</i> out of Africa. Evolutionary Anthropology, 2015, 24, 149-164.	3.4	263
22	Reconstructing regional population fluctuations in the European Neolithic using radiocarbon dates: a new case-study using an improved method. Journal of Archaeological Science, 2014, 52, 549-557.	2.4	262
23	A worldwide correlation of lactase persistence phenotype and genotypes. BMC Evolutionary Biology, 2010, 10, 36.	3.2	258
24	Ancient Hybridization and an Irish Origin for the Modern Polar Bear Matriline. Current Biology, 2011, 21, 1251-1258.	3.9	257
25	Modeling Recent Human Evolution in Mice by Expression of a Selected EDAR Variant. Cell, 2013, 152, 691-702.	28.9	250
26	A novel polymorphism associated with lactose tolerance in Africa: multiple causes for lactase persistence?. Human Genetics, 2007, 120, 779-788.	3.8	247
27	Direct evidence for positive selection of skin, hair, and eye pigmentation in Europeans during the last 5,000 y. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 4832-4837.	7.1	240
28	Antibodies to conformational epitopes of soluble liver antigen define a severe form of autoimmune liver disease. Hepatology, 2002, 35, 658-664.	7.3	236
29	Early Neolithic genomes from the eastern Fertile Crescent. Science, 2016, 353, 499-503.	12.6	230
30	Ancient DNA Reveals Lack of Continuity between Neolithic Hunter-Gatherers and Contemporary Scandinavians. Current Biology, 2009, 19, 1758-1762.	3.9	217
31	Mitochondrial DNA analysis shows a Near Eastern Neolithic origin for domestic cattle and no indication of domestication of European aurochs. Proceedings of the Royal Society B: Biological Sciences, 2007, 274, 1377-1385.	2.6	209
32	Radiocarbon evidence indicates that migrants introduced farming to Britain. Journal of Archaeological Science, 2010, 37, 866-870.	2.4	199
33	The T Allele of a Single-Nucleotide Polymorphism 13.9 kb Upstream of the Lactase Gene (LCT) (C _{13.9} kbT) Does Not Predict or Cause the Lactase-Persistence Phenotype in Africans. American Journal of Human Genetics, 2004, 74, 1102-1110.	6.2	196
34	Evidence for a Common Origin of Blacksmiths and Cultivators in the Ethiopian Ari within the Last 4500 Years: Lessons for Clustering-Based Inference. PLoS Genetics, 2015, 11, e1005397.	3.5	194
35	Making evolutionary biology a basic science for medicine. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 1800-1807.	7.1	189
36	A Y Chromosome Census of the British Isles. Current Biology, 2003, 13, 979-984.	3.9	185

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37	Genetic evidence for different male and female roles during cultural transitions in the British Isles. Proceedings of the National Academy of Sciences of the United States of America, 2001, 98, 5078-5083.	7.1	182
38	Resolution of chronic hepatitis B and anti-HBs seroconversion in humans by adoptive transfer of immunity to hepatitis B core antigen. Gastroenterology, 2002, 122, 614-624.	1.3	180
39	Y-Chromosome Evidence for Differing Ancient Demographic Histories in the Americas. American Journal of Human Genetics, 2003, 73, 524-539.	6.2	180
40	Ethiopian Genetic Diversity Reveals Linguistic Stratification and Complex Influences on the Ethiopian Gene Pool. American Journal of Human Genetics, 2012, 91, 83-96.	6.2	177
41	Y Chromosomes Traveling South: The Cohen Modal Haplotype and the Origins of the Lembaâ€”the â€œBlack Jews of Southern Africaâ€”. American Journal of Human Genetics, 2000, 66, 674-686.	6.2	174
42	Origins of Old Testament priests. Nature, 1998, 394, 138-140.	27.8	170
43	The Importance of Dietary Carbohydrate in Human Evolution. Quarterly Review of Biology, 2015, 90, 251-268.	0.1	168
44	2000 Years of Parallel Societies in Stone Age Central Europe. Science, 2013, 342, 479-481.	12.6	165
45	Identification of the remains of King Richard III. Nature Communications, 2014, 5, 5631.	12.8	163
46	Ancient genomes indicate population replacement in Early Neolithic Britain. Nature Ecology and Evolution, 2019, 3, 765-771.	7.8	156
47	<i>AIP</i> Mutation in Pituitary Adenomas in the 18th Century and Today. New England Journal of Medicine, 2011, 364, 43-50.	27.0	151
48	Y Chromosome Evidence for Anglo-Saxon Mass Migration. Molecular Biology and Evolution, 2002, 19, 1008-1021.	8.9	148
49	Modern Taurine Cattle Descended from Small Number of Near-Eastern Founders. Molecular Biology and Evolution, 2012, 29, 2101-2104.	8.9	131
50	Staying out in the cold: glacial refugia and mitochondrial DNA phylogeography in ancient European brown bears. Molecular Ecology, 2007, 16, 5140-5148.	3.9	130
51	Founding Mothers of Jewish Communities: Geographically Separated Jewish Groups Were Independently Founded by Very Few Female Ancestors. American Journal of Human Genetics, 2002, 70, 1411-1420.	6.2	126
52	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.	6.2	124
53	Understanding cumulative cultural evolution. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, E6724-E6725.	7.1	124
54	Metallothionein genes from the flowering plant <i>Mimulus guttatus</i> . FEBS Letters, 1990, 260, 277-280.	2.8	110

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55	Genetic Structure and Extinction of the Woolly Mammoth, <i>Mammuthus primigenius</i> . <i>Current Biology</i> , 2007, 17, 1072-1075.	3.9	109
56	ANCIENT URBANIZATION PREDICTS GENETIC RESISTANCE TO TUBERCULOSIS. <i>Evolution; International Journal of Organic Evolution</i> , 2011, 65, 842-848.	2.3	108
57	Obesity, starch digestion and amylase: association between copy number variants at human salivary (AMY1) and pancreatic (AMY2) amylase genes. <i>Human Molecular Genetics</i> , 2015, 24, 3472-3480.	2.9	105
58	The genetic history of Europeans. <i>Trends in Genetics</i> , 2012, 28, 496-505.	6.7	102
59	Excavating Past Population Structures by Surname-Based Sampling: The Genetic Legacy of the Vikings in Northwest England. <i>Molecular Biology and Evolution</i> , 2008, 25, 301-309.	8.9	101
60	Genomic signals of migration and continuity in Britain before the Anglo-Saxons. <i>Nature Communications</i> , 2016, 7, 10326.	12.8	100
61	The evolution of lactase persistence in Europe. A synthesis of archaeological and genetic evidence. <i>International Dairy Journal</i> , 2012, 22, 88-97.	3.0	97
62	Storytelling and story testing in domestication. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 6159-6164.	7.1	96
63	DNA from ancient mammoth bones. <i>Nature</i> , 1994, 370, 333-334.	27.8	92
64	Characterization of hunter-gatherer networks and implications for cumulative culture. <i>Nature Human Behaviour</i> , 2017, 1, .	12.0	91
65	Multiple Origins of Ashkenazi Levites: Y Chromosome Evidence for Both Near Eastern and European Ancestries. <i>American Journal of Human Genetics</i> , 2003, 73, 768-779.	6.2	90
66	Multiple Rare Variants as a Cause of a Common Phenotype: Several Different Lactase Persistence Associated Alleles in a Single Ethnic Group. <i>Journal of Molecular Evolution</i> , 2009, 69, 579-588.	1.8	89
67	The phylogenetic position of the "giant deer" <i>Megaloceros giganteus</i> . <i>Nature</i> , 2005, 438, 850-853.	27.8	88
68	Surprising migration and population size dynamics in ancient Iberian brown bears (<i>Ursus arctos</i>). <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 5123-5128.	7.1	86
69	Ancient West African foragers in the context of African population history. <i>Nature</i> , 2020, 577, 665-670.	27.8	86
70	Large-scale migration into Britain during the Middle to Late Bronze Age. <i>Nature</i> , 2022, 601, 588-594.	27.8	86
71	High throughput analysis of 10 microsatellite and 11 diallelic polymorphisms on the human Y-chromosome. <i>Human Genetics</i> , 1999, 105, 577-581.	3.8	85
72	Oral microbiomes from hunter-gatherers and traditional farmers reveal shifts in commensal balance and pathogen load linked to diet. <i>Molecular Ecology</i> , 2018, 27, 182-195.	3.9	85

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73	The peopling of Europe and the cautionary tale of Y chromosome lineage R-M269. Proceedings of the Royal Society B: Biological Sciences, 2012, 279, 884-892.	2.6	84
74	World-wide distributions of lactase persistence alleles and the complex effects of recombination and selection. Human Genetics, 2017, 136, 1445-1453.	3.8	81
75	The Arrival of Siberian Ancestry Connecting the Eastern Baltic to Uralic Speakers further East. Current Biology, 2019, 29, 1701-1711.e16.	3.9	80
76	Evidence for an apartheid-like social structure in early Anglo-Saxon England. Proceedings of the Royal Society B: Biological Sciences, 2006, 273, 2651-2657.	2.6	79
77	Normophosphatemic Familial Tumoral Calcinosis Is Caused by Deleterious Mutations in SAMD9, Encoding a TNF- α Responsive Protein. Journal of Investigative Dermatology, 2008, 128, 1423-1429.	0.7	76
78	Key Residues of a Major Cytochrome P4502D6 Epitope Are Located on the Surface of the Molecule. Journal of Immunology, 2002, 169, 277-285.	0.8	74
79	High-resolution Y chromosome haplotypes of Israeli and Palestinian Arabs reveal geographic substructure and substantial overlap with haplotypes of Jews. Human Genetics, 2000, 107, 630-641.	3.8	69
80	Beyond multiregional and simple out-of-Africa models of human evolution. Nature Ecology and Evolution, 2019, 3, 1370-1372.	7.8	68
81	Herders of Indian and European Cattle Share Their Predominant Allele for Lactase Persistence. Molecular Biology and Evolution, 2012, 29, 249-260.	8.9	67
82	Genomic variations in the hepatitis B core gene: A possible factor influencing response to interferon alfa treatment. Gastroenterology, 1995, 108, 505-514.	1.3	65
83	Evolution of a Length Polymorphism in the Human PER3 Gene, a Component of the Circadian System. Journal of Biological Rhythms, 2005, 20, 490-499.	2.6	64
84	Inferring Allele Frequency Trajectories from Ancient DNA Indicates That Selection on a Chicken Gene Coincided with Changes in Medieval Husbandry Practices. Molecular Biology and Evolution, 2017, 34, 1981-1990.	8.9	63
85	Armenian Y chromosome haplotypes reveal strong regional structure within a single ethno-national group. Human Genetics, 2001, 109, 659-674.	3.8	58
86	Little genetic differentiation as assessed by uniparental markers in the presence of substantial language variation in peoples of the Cross River region of Nigeria. BMC Evolutionary Biology, 2010, 10, 92.	3.2	57
87	Population Structure in the Mediterranean Basin: A Y Chromosome Perspective. Annals of Human Genetics, 2006, 70, 207-225.	0.8	56
88	Low Prevalence of Lactase Persistence in Bronze Age Europe Indicates Ongoing Strong Selection over the Last 3,000 Years. Current Biology, 2020, 30, 4307-4315.e13.	3.9	54
89	Prevalence of Clinically Relevant UGT1A Alleles and Haplotypes in African Populations. Annals of Human Genetics, 2011, 75, 236-246.	0.8	53
90	Case against subclassification of type II autoimmune chronic active hepatitis. Lancet, The, 1993, 341, 60.	13.7	52

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91	Direct Estimates of Natural Selection in Iberia Indicate Calcium Absorption Was Not the Only Driver of Lactase Persistence in Europe. <i>Molecular Biology and Evolution</i> , 2014, 31, 975-983.	8.9	52
92	Hepatitis B virus variants with core gene deletions in the evolution of chronic hepatitis B infection. <i>Gastroenterology</i> , 1996, 111, 183-192.	1.3	49
93	Molecular diversity and population structure at the Cytochrome P450 3A5 gene in Africa. <i>BMC Genetics</i> , 2013, 14, 34.	2.7	49
94	Disentangling Immediate Adaptive Introgression from Selection on Standing Introgressed Variation in Humans. <i>Molecular Biology and Evolution</i> , 2018, 35, 623-630.	8.9	46
95	A 3,000-year-old Egyptian emmer wheat genome reveals dispersal and domestication history. <i>Nature Plants</i> , 2019, 5, 1120-1128.	9.3	46
96	Molecular and morphological evidence on the phylogeny of the Elephantidae. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2000, 267, 2493-2500.	2.6	45
97	Evaluating bacterial pathogen DNA preservation in museum osteological collections. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2006, 273, 645-653.	2.6	44
98	The potentially deleterious functional variant flavin-containing monooxygenase 2*1 is at high frequency throughout sub-Saharan Africa. <i>Pharmacogenetics and Genomics</i> , 2008, 18, 877-886.	1.5	43
99	Increased prevalence of M694V in patients with ankylosing spondylitis: Additional evidence for a link with familial mediterranean fever. <i>Arthritis and Rheumatism</i> , 2010, 62, 3059-3063.	6.7	43
100	Estimating mobility using sparse data: Application to human genetic variation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 12213-12218.	7.1	37
101	XAF1 as a modifier of p53 function and cancer susceptibility. <i>Science Advances</i> , 2020, 6, eaba3231.	10.3	37
102	A Rare Deep-Rooting DO African Y-Chromosomal Haplogroup and Its Implications for the Expansion of Modern Humans Out of Africa. <i>Genetics</i> , 2019, 212, 1421-1428.	2.9	35
103	High throughput analysis of 10 microsatellite and 11 diallelic polymorphisms on the human Y-chromosome. <i>Human Genetics</i> , 1999, 105, 577-581.	3.8	34
104	Population history of the Hispaniolan hutia <i>Plagiodontia aedium</i> (Rodentia: Capromyidae): testing the model of ancient differentiation on a geotectonically complex Caribbean island. <i>Molecular Ecology</i> , 2012, 21, 2239-2253.	3.9	34
105	How long have adult humans been consuming milk?. <i>IUBMB Life</i> , 2013, 65, 983-990.	3.4	34
106	Accurate age estimation in small-scale societies. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 8205-8210.	7.1	33
107	Statistically robust representation and comparison of mortality profiles in archaeozoology. <i>Journal of Archaeological Science</i> , 2016, 71, 24-32.	2.4	31
108	Palaeoecological and genetic evidence for Neanderthal power locomotion as an adaptation to a woodland environment. <i>Quaternary Science Reviews</i> , 2019, 217, 310-315.	3.0	31

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109	Diet and the frequency of the alanine:glyoxylate aminotransferase Pro11Leu polymorphism in different human populations. <i>Human Genetics</i> , 2004, 115, 504-509.	3.8	30
110	50,000 years of genetic uniformity in the critically endangered Iberian lynx. <i>Molecular Ecology</i> , 2011, 20, 3785-3795.	3.9	30
111	Rare Deep-Rooting Y Chromosome Lineages in Humans: Lessons for Phylogeography. <i>Genetics</i> , 2003, 165, 229-234.	2.9	29
112	Ribeiro's typology, genomes, and Spanish colonialism, as viewed from Gran Canaria and Colombia. <i>Genetics and Molecular Biology</i> , 2004, 27, 01-08.	1.3	28
113	CYP1A2 is more variable than previously thought: a genomic biography of the gene behind the human drug-metabolizing enzyme. <i>Pharmacogenetics and Genomics</i> , 2010, 20, 647-664.	1.5	27
114	The Greeks in the West: genetic signatures of the Hellenic colonisation in southern Italy and Sicily. <i>European Journal of Human Genetics</i> , 2016, 24, 429-436.	2.8	26
115	Increased Population Risk of AIP-Related Acromegaly and Gigantism in Ireland. <i>Human Mutation</i> , 2017, 38, 78-85.	2.5	25
116	Evidence of the interplay of genetics and culture in Ethiopia. <i>Nature Communications</i> , 2021, 12, 3581.	12.8	25
117	Molecular instability in the COL11A2 tRNA ^{Lys} intergenic region of the human mitochondrial genome: multiple origins of the 9bp deletion and heteroplasmy for expanded repeats. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 1998, 353, 955-965.	4.0	23
118	The Use of Y-Chromosomal DNA Variation to Investigate Population History. , 1999, , 91-101.		23
119	From a dry bone to a genetic portrait: A case study of sickle cell anemia. , 2000, 111, 153-163.		23
120	Integration versus apartheid in post-Roman Britain: a response to Pattison. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2008, 275, 2419-2421.	2.6	23
121	Transition to farming more likely for small, conservative groups with property rights, but increased productivity is not essential. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 14218-14223.	7.1	23
122	Genetic evidence for a western Chinese origin of broomcorn millet (<i>Panicum miliaceum</i>). <i>Holocene</i> , 2018, 28, 1968-1978.	1.7	23
123	Directly modelling population dynamics in the South American Arid Diagonal using ¹⁴ C dates. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2021, 376, 20190723.	4.0	23
124	An Analysis of Consanguinity and Social Structure Within the UK Asian Population Using Microsatellite Data. <i>Annals of Human Genetics</i> , 2003, 67, 525-537.	0.8	22
125	Genetic diversity of CHC22 clathrin impacts its function in glucose metabolism. <i>ELife</i> , 2019, 8, .	6.0	22
126	An assessment of the long-term preservation of the DNA of a bacterial pathogen in ethanol-preserved archival material. <i>Journal of Pathology</i> , 2000, 192, 554-559.	4.5	21

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127	Molecular phylogeny of genus <i>Guizotia</i> (Asteraceae) using DNA sequences derived from ITS. <i>Genetic Resources and Crop Evolution</i> , 2007, 54, 1419-1427.	1.6	20
128	Reduced intensity of bone fat exploitation correlates with increased potential access to dairy fats in early Neolithic Europe. <i>Journal of Archaeological Science</i> , 2018, 94, 60-69.	2.4	20
129	The Genetic Legacy of Zoroastrianism in Iran and India: Insights into Population Structure, Gene Flow, and Selection. <i>American Journal of Human Genetics</i> , 2017, 101, 353-368.	6.2	19
130	Diversity of lactase persistence in African milk drinkers. <i>Human Genetics</i> , 2015, 134, 917-925.	3.8	18
131	New genetic evidence supports isolation and drift in the Ladin communities of the South Tyrolean Alps but not an ancient origin in the Middle East. <i>European Journal of Human Genetics</i> , 2008, 16, 124-134.	2.8	17
132	Independent evolutionary histories in allopatric populations of a threatened Caribbean land mammal. <i>Diversity and Distributions</i> , 2016, 22, 589-602.	4.1	17
133	Synchronous diversification of Sulawesi's iconic artiodactyls driven by recent geological events. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2018, 285, 20172566.	2.6	17
134	Palenque de San Basilio in Colombia: genetic data support an oral history of a paternal ancestry in Congo. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2016, 283, 20152980.	2.6	14
135	Three Reportedly Unrelated Families With Liddle Syndrome Inherited From a Common Ancestor. <i>Hypertension</i> , 2018, 71, 273-279.	2.7	14
136	Simulating Geographical Variation in Material Culture: Were Early Modern Humans in Europe Ethnically Structured?. , 2015, , 103-120.		13
137	Summary: The Science of Genealogy by Genetics. <i>Developing World Bioethics</i> , 2003, 3, 103-108.	0.9	12
138	In-frame seven amino-acid duplication in AIP arose over the last 3000 years, disrupts protein interaction and stability and is associated with gigantism. <i>European Journal of Endocrinology</i> , 2017, 177, 257-266.	3.7	12
139	Sequencing of cDNA using anchored oligo dT primers. <i>Nucleic Acids Research</i> , 1993, 21, 3915-3916.	14.5	11
140	Y chromosome haplotypes and testicular cancer in the English population. <i>Journal of Medical Genetics</i> , 2003, 40, 20e-20.	3.2	11
141	Assessing the effects of conservation treatments on short sequences of DNA in vitro. <i>Journal of Archaeological Science</i> , 2010, 37, 2831-2841.	2.4	11
142	Evaluating demographic models for goat domestication using mtDNA sequences. <i>Anthropozoologica</i> , 2012, 47, 64-76.	0.5	11
143	Preparation of Bone Samples for DNA Extraction: A Nuts and Bolts Approach. <i>BioTechniques</i> , 1997, 22, 402-402.	1.8	9
144	High-throughput analysis of informative CYP2D6 compound haplotypes. <i>Genomics</i> , 2003, 81, 166-174.	2.9	9

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145	The flickering genes of the last mammoths. <i>Molecular Ecology</i> , 2012, 21, 3379-3381.	3.9	9
146	Food Income and the Evolution of Forager Mobility. <i>Scientific Reports</i> , 2019, 9, 5438.	3.3	9
147	Genetic legacy of state centralization in the Kuba Kingdom of the Democratic Republic of the Congo. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 593-598.	7.1	9
148	Human origins in Southern African palaeo-wetlands? Strong claims from weak evidence. <i>Journal of Archaeological Science</i> , 2021, 130, 105374.	2.4	9
149	A method for avoiding mis-priming when sequencing with Dynabeads. <i>Nucleic Acids Research</i> , 1994, 22, 3243-3244.	14.5	8
150	Sex-specific Genetic Data Support One of Two Alternative Versions of the Foundation of the Ruling Dynasty of the Nso in Cameroon. <i>Current Anthropology</i> , 2008, 49, 707-714.	1.6	8
151	Pitfalls of the Geographic Population Structure (GPS) Approach Applied to Human Genetic History: A Case Study of Ashkenazi Jews. <i>Genome Biology and Evolution</i> , 2016, 8, 2259-2265.	2.5	7
152	Investigating mitochondrial DNA relationships in Neolithic Western Europe through serial coalescent simulations. <i>European Journal of Human Genetics</i> , 2017, 25, 388-392.	2.8	7
153	Demography and Variation in the Accumulation of Culturally Inherited Skills. , 2009, , 137-160.		7
154	<i>Candida lusitanae</i> septicemia: Successful combination therapy. <i>Clinical Microbiology Newsletter</i> , 1985, 7, 142-143.	0.7	6
155	Detecting Gene Duplications in the Human Lineage. <i>Annals of Human Genetics</i> , 2010, 74, 555-565.	0.8	6
156	Long-term DNA survival in ethanol-preserved archival material. <i>Annals of the Royal College of Surgeons of England</i> , 2001, 83, 283-4.	0.6	5
157	Modelling caprine age-at-death profiles using the Gamma distribution. <i>Journal of Archaeological Science</i> , 2018, 99, 19-26.	2.4	4
158	The evolution of lactose digestion. , 2019, , 1-48.		4
159	Lactose intolerance and other related food sensitivities. , 2019, , 113-153.		4
160	The Use of PCR for Differential Screening of cDNA Libraries. , 1997, 67, 405-418.		3
161	Mitochondrial DNA and IQ in Europe. <i>Intelligence</i> , 1998, 26, 167-173.	3.0	3
162	“Ava”: a Beaker-associated woman from a cist at Achavanich, Highland, and the story of her (re-)discovery and subsequent study. <i>Proceedings of the Society of Antiquaries of Scotland</i> , 0, 147, 73-118.	0.0	3

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163	TT Virus Infection in Patients with Primary Hypogammaglobulinaemia: Natural History and Relationship to Liver Disease in the Immunocompromised Host. <i>Scandinavian Journal of Gastroenterology</i> , 2001, 36, 987-993.	1.5	2
164	Y chromosomes of self-identified Syeds from the Indian subcontinent show evidence of elevated Arab ancestry but not of a recent common patrilineal origin. <i>Archaeological and Anthropological Sciences</i> , 2010, 2, 217-224.	1.8	2
165	Genetic and isotopic analysis and the UK Border Agency. <i>Significance</i> , 2010, 7, 58-61.	0.4	2
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