

# David Reich

## List of Publications by Citations

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168  
papers

38,014  
citations

69  
h-index

187  
g-index

187  
ext. papers

48,426  
ext. citations

22.7  
avg, IF

7.02  
L-index

#	Paper	IF	Citations
168	Principal components analysis corrects for stratification in genome-wide association studies. <i>Nature Genetics</i> , <b>2006</b> , 38, 904-9	36.3	7055
167	Population structure and eigenanalysis. <i>PLoS Genetics</i> , <b>2006</b> , 2, e190	6	3021
166	A draft sequence of the Neandertal genome. <i>Science</i> , <b>2010</b> , 328, 710-722	33.3	2599
165	The complete genome sequence of a Neanderthal from the Altai Mountains. <i>Nature</i> , <b>2014</b> , 505, 43-9	50.4	1339
164	A high-coverage genome sequence from an archaic Denisovan individual. <i>Science</i> , <b>2012</b> , 338, 222-6	33.3	1276
163	Ancient admixture in human history. <i>Genetics</i> , <b>2012</b> , 192, 1065-93	4	1212
162	Genetic history of an archaic hominin group from Denisova Cave in Siberia. <i>Nature</i> , <b>2010</b> , 468, 1053-60	50.4	1169
161	Reconstructing Indian population history. <i>Nature</i> , <b>2009</b> , 461, 489-94	50.4	1075
160	Massive migration from the steppe was a source for Indo-European languages in Europe. <i>Nature</i> , <b>2015</b> , 522, 207-11	50.4	968
159	Ancient human genomes suggest three ancestral populations for present-day Europeans. <i>Nature</i> , <b>2014</b> , 513, 409-13	50.4	812
158	Genome-wide patterns of selection in 230 ancient Eurasians. <i>Nature</i> , <b>2015</b> , 528, 499-503	50.4	774
157	The Simons Genome Diversity Project: 300 genomes from 142 diverse populations. <i>Nature</i> , <b>2016</b> , 538, 201-206	50.4	759
156	Testing for ancient admixture between closely related populations. <i>Molecular Biology and Evolution</i> , <b>2011</b> , 28, 2239-52	8.3	689
155	Cost-effective, high-throughput DNA sequencing libraries for multiplexed target capture. <i>Genome Research</i> , <b>2012</b> , 22, 939-46	9.7	637
154	Genome sequence of a 45,000-year-old modern human from western Siberia. <i>Nature</i> , <b>2014</b> , 514, 445-9	50.4	635
153	The genomic landscape of Neanderthal ancestry in present-day humans. <i>Nature</i> , <b>2014</b> , 507, 354-7	50.4	615
152	Reconstructing Native American population history. <i>Nature</i> , <b>2012</b> , 488, 370-4	50.4	498

151	Genomic insights into the origin of farming in the ancient Near East. <i>Nature</i> , <b>2016</b> , 536, 419-24	50.4	485
150	The genetic history of Ice Age Europe. <i>Nature</i> , <b>2016</b> , 534, 200-5	50.4	473
149	An early modern human from Romania with a recent Neanderthal ancestor. <i>Nature</i> , <b>2015</b> , 524, 216-9	50.4	446
148	Denisova admixture and the first modern human dispersals into Southeast Asia and Oceania. <i>American Journal of Human Genetics</i> , <b>2011</b> , 89, 516-28	11	390
147	A revised timescale for human evolution based on ancient mitochondrial genomes. <i>Current Biology</i> , <b>2013</b> , 23, 553-559	6.3	387
146	Methods for high-density admixture mapping of disease genes. <i>American Journal of Human Genetics</i> , <b>2004</b> , 74, 979-1000	11	386
145	The genetic ancestry of African Americans, Latinos, and European Americans across the United States. <i>American Journal of Human Genetics</i> , <b>2015</b> , 96, 37-53	11	329
144	A high-coverage Neanderthal genome from Vindija Cave in Croatia. <i>Science</i> , <b>2017</b> , 358, 655-658	33.3	312
143	The Beaker phenomenon and the genomic transformation of northwest Europe. <i>Nature</i> , <b>2018</b> , 555, 190-194	30.4	293
142	Inferring admixture histories of human populations using linkage disequilibrium. <i>Genetics</i> , <b>2013</b> , 193, 1233-54	4	293
141	The genomic history of southeastern Europe. <i>Nature</i> , <b>2018</b> , 555, 197-203	50.4	287
140	The Combined Landscape of Denisovan and Neanderthal Ancestry in Present-Day Humans. <i>Current Biology</i> , <b>2016</b> , 26, 1241-7	6.3	261
139	Reduced neutrophil count in people of African descent is due to a regulatory variant in the Duffy antigen receptor for chemokines gene. <i>PLoS Genetics</i> , <b>2009</b> , 5, e1000360	6	251
138	Partial uracil-DNA-glycosylase treatment for screening of ancient DNA. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , <b>2015</b> , 370, 20130624	5.8	249
137	The landscape of recombination in African Americans. <i>Nature</i> , <b>2011</b> , 476, 170-5	50.4	243
136	Gibbon genome and the fast karyotype evolution of small apes. <i>Nature</i> , <b>2014</b> , 513, 195-201	50.4	241
135	Ancient DNA reveals key stages in the formation of central European mitochondrial genetic diversity. <i>Science</i> , <b>2013</b> , 342, 257-61	33.3	237
134	Global diversity, population stratification, and selection of human copy-number variation. <i>Science</i> , <b>2015</b> , 349, aab3761	33.3	224

133	Genetic evidence for two founding populations of the Americas. <i>Nature</i> , <b>2015</b> , 525, 104-8	50.4	220
132	A whole-genome admixture scan finds a candidate locus for multiple sclerosis susceptibility. <i>Nature Genetics</i> , <b>2005</b> , 37, 1113-8	36.3	220
131	Measurement of the human allele frequency spectrum demonstrates greater genetic drift in East Asians than in Europeans. <i>Nature Genetics</i> , <b>2007</b> , 39, 1251-5	36.3	212
130	Ancient mitochondrial DNA provides high-resolution time scale of the peopling of the Americas. <i>Science Advances</i> , <b>2016</b> , 2, e1501385	14.3	211
129	Reconstructing Prehistoric African Population Structure. <i>Cell</i> , <b>2017</b> , 171, 59-71.e21	56.2	201
128	Insights into human genetic variation and population history from 929 diverse genomes. <i>Science</i> , <b>2020</b> , 367,	33.3	196
127	The formation of human populations in South and Central Asia. <i>Science</i> , <b>2019</b> , 365,	33.3	195
126	Parallel palaeogenomic transects reveal complex genetic history of early European farmers. <i>Nature</i> , <b>2017</b> , 551, 368-372	50.4	194
125	The genetic prehistory of southern Africa. <i>Nature Communications</i> , <b>2012</b> , 3, 1143	17.4	193
124	The genomic history of the Iberian Peninsula over the past 8000 years. <i>Science</i> , <b>2019</b> , 363, 1230-1234	33.3	186
123	Genomic insights into the peopling of the Southwest Pacific. <i>Nature</i> , <b>2016</b> , 538, 510-513	50.4	180
122	Complete genomes reveal signatures of demographic and genetic declines in the woolly mammoth. <i>Current Biology</i> , <b>2015</b> , 25, 1395-400	6.3	177
121	Genetic evidence for recent population mixture in India. <i>American Journal of Human Genetics</i> , <b>2013</b> , 93, 422-38	11	177
120	Ancient west Eurasian ancestry in southern and eastern Africa. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2014</b> , 111, 2632-7	11.5	176
119	The history of African gene flow into Southern Europeans, Levantines, and Jews. <i>PLoS Genetics</i> , <b>2011</b> , 7, e1001373	6	175
118	Polygenic adaptation on height is overestimated due to uncorrected stratification in genome-wide association studies. <i>ELife</i> , <b>2019</b> , 8,	8.9	166
117	Toward a new history and geography of human genes informed by ancient DNA. <i>Trends in Genetics</i> , <b>2014</b> , 30, 377-89	8.5	163
116	Ancient genomes document multiple waves of migration in Southeast Asian prehistory. <i>Science</i> , <b>2018</b> , 361, 92-95	33.3	149

115	Admixture mapping of an allele affecting interleukin 6 soluble receptor and interleukin 6 levels. <i>American Journal of Human Genetics</i> , <b>2007</b> , 80, 716-26	11	145
114	Reconstructing the Deep Population History of Central and South America. <i>Cell</i> , <b>2018</b> , 175, 1185-1197.e23	36.2	143
113	Reconstructing the genetic history of late Neanderthals. <i>Nature</i> , <b>2018</b> , 555, 652-656	50.4	138
112	Archaeogenomic evidence reveals prehistoric matrilineal dynasty. <i>Nature Communications</i> , <b>2017</b> , 8, 14115	17.4	134
111	No evidence that selection has been less effective at removing deleterious mutations in Europeans than in Africans. <i>Nature Genetics</i> , <b>2015</b> , 47, 126-31	36.3	129
110	Reconstructing Austronesian population history in Island Southeast Asia. <i>Nature Communications</i> , <b>2014</b> , 5, 4689	17.4	116
109	Genetic origins of the Minoans and Mycenaeans. <i>Nature</i> , <b>2017</b> , 548, 214-218	50.4	108
108	The contribution of rare variation to prostate cancer heritability. <i>Nature Genetics</i> , <b>2016</b> , 48, 30-5	36.3	106
107	The genetic prehistory of the Baltic Sea region. <i>Nature Communications</i> , <b>2018</b> , 9, 442	17.4	96
106	Ancient DNA indicates human population shifts and admixture in northern and southern China. <i>Science</i> , <b>2020</b> , 369, 282-288	33.3	90
105	A comprehensive genomic history of extinct and living elephants. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2018</b> , 115, E2566-E2574	11.5	86
104	Ancient genomes indicate population replacement in Early Neolithic Britain. <i>Nature Ecology and Evolution</i> , <b>2019</b> , 3, 765-771	12.3	82
103	The promise of discovering population-specific disease-associated genes in South Asia. <i>Nature Genetics</i> , <b>2017</b> , 49, 1403-1407	36.3	79
102	Efficient moment-based inference of admixture parameters and sources of gene flow. <i>Molecular Biology and Evolution</i> , <b>2013</b> , 30, 1788-802	8.3	78
101	A genetic method for dating ancient genomes provides a direct estimate of human generation interval in the last 45,000 years. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2016</b> , 113, 5652-7	11.5	75
100	Will admixture mapping work to find disease genes?. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , <b>2005</b> , 360, 1605-7	5.8	70
99	The genetic history of admixture across inner Eurasia. <i>Nature Ecology and Evolution</i> , <b>2019</b> , 3, 966-976	12.3	69
98	Current evidence allows multiple models for the peopling of the Americas. <i>Science Advances</i> , <b>2018</b> , 4, eaat5473	14.3	63

97	A genomic view of the peopling of the Americas. <i>Current Opinion in Genetics and Development</i> , <b>2016</b> , 41, 27-35	4.9	63
96	Genomic insights into the formation of human populations in East Asia. <i>Nature</i> , <b>2021</b> , 591, 413-419	50.4	62
95	Ancestry and demography and descendants of Iron Age nomads of the Eurasian Steppe. <i>Nature Communications</i> , <b>2017</b> , 8, 14615	17.4	60
94	Non-crossover gene conversions show strong GC bias and unexpected clustering in humans. <i>ELife</i> , <b>2015</b> , 4,	8.9	57
93	Differences in the rare variant spectrum among human populations. <i>PLoS Genetics</i> , <b>2017</b> , 13, e1006581	6	56
92	Population Turnover in Remote Oceania Shortly after Initial Settlement. <i>Current Biology</i> , <b>2018</b> , 28, 1157-1165	13.65	55
91	Ancient human genome-wide data from a 3000-year interval in the Caucasus corresponds with eco-geographic regions. <i>Nature Communications</i> , <b>2019</b> , 10, 590	17.4	55
90	Ancient DNA reveals a multistep spread of the first herders into sub-Saharan Africa. <i>Science</i> , <b>2019</b> , 365,	33.3	53
89	Palaeo-Eskimo genetic ancestry and the peopling of Chukotka and North America. <i>Nature</i> , <b>2019</b> , 570, 236-240	50.4	53
88	African evolutionary history inferred from whole genome sequence data of 44 indigenous African populations. <i>Genome Biology</i> , <b>2019</b> , 20, 82	18.3	53
87	Dominance of Deleterious Alleles Controls the Response to a Population Bottleneck. <i>PLoS Genetics</i> , <b>2015</b> , 11, e1005436	6	53
86	Extreme selective sweeps independently targeted the X chromosomes of the great apes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2015</b> , 112, 6413-8	11.5	52
85	The spread of steppe and Iranian-related ancestry in the islands of the western Mediterranean. <i>Nature Ecology and Evolution</i> , <b>2020</b> , 4, 334-345	12.3	48
84	Ancient West African foragers in the context of African population history. <i>Nature</i> , <b>2020</b> , 577, 665-670	50.4	47
83	Genome-wide scan of 29,141 African Americans finds no evidence of directional selection since admixture. <i>American Journal of Human Genetics</i> , <b>2014</b> , 95, 437-44	11	46
82	Leveraging population admixture to characterize the heritability of complex traits. <i>Nature Genetics</i> , <b>2014</b> , 46, 1356-62	36.3	45
81	Genetic history from the Middle Neolithic to present on the Mediterranean island of Sardinia. <i>Nature Communications</i> , <b>2020</b> , 11, 939	17.4	42
80	A Working Model of the Deep Relationships of Diverse Modern Human Genetic Lineages Outside of Africa. <i>Molecular Biology and Evolution</i> , <b>2017</b> , 34, 889-902	8.3	39

79	Natural hybridization reveals incompatible alleles that cause melanoma in swordtail fish. <i>Science</i> , <b>2020</b> , 368, 731-736	33.3	38
78	Palaeogenomes of Eurasian straight-tusked elephants challenge the current view of elephant evolution. <i>ELife</i> , <b>2017</b> , 6,	8.9	37
77	Ancient DNA from Chalcolithic Israel reveals the role of population mixture in cultural transformation. <i>Nature Communications</i> , <b>2018</b> , 9, 3336	17.4	37
76	Interpreting short tandem repeat variations in humans using mutational constraint. <i>Nature Genetics</i> , <b>2017</b> , 49, 1495-1501	36.3	37
75	Calibrating the Human Mutation Rate via Ancestral Recombination Density in Diploid Genomes. <i>PLoS Genetics</i> , <b>2015</b> , 11, e1005550	6	37
74	Reconciling material cultures in archaeology with genetic data: The nomenclature of clusters emerging from archaeogenomic analysis. <i>Scientific Reports</i> , <b>2018</b> , 8, 13003	4.9	36
73	A Re-Appraisal of the Early Andean Human Remains from Lauricocha in Peru. <i>PLoS ONE</i> , <b>2015</b> , 10, e0127141	3.4	34
72	A Paleogenomic Reconstruction of the Deep Population History of the Andes. <i>Cell</i> , <b>2020</b> , 181, 1131-1145	62.1	33
71	A genetic history of the pre-contact Caribbean. <i>Nature</i> , <b>2021</b> , 590, 103-110	50.4	32
70	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. <i>Genes</i> , <b>2018</b> , 9,	4.2	30
69	An Ancient Harappan Genome Lacks Ancestry from Steppe Pastoralists or Iranian Farmers. <i>Cell</i> , <b>2019</b> , 179, 729-735.e10	56.2	28
68	Differential DNA methylation of vocal and facial anatomy genes in modern humans. <i>Nature Communications</i> , <b>2020</b> , 11, 1189	17.4	27
67	The Promise of Paleogenomics Beyond Our Own Species. <i>Trends in Genetics</i> , <b>2019</b> , 35, 319-329	8.5	22
66	Arrival routes of first Americans uncertain. <i>Science</i> , <b>2018</b> , 359, 1224-1225	33.3	22
65	Evaluating the contribution of rare variants to type 2 diabetes and related traits using pedigrees. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2018</b> , 115, 379-384	11.5	21
64	Genome diversity in the Neolithic Globular Amphorae culture and the spread of Indo-European languages. <i>Proceedings of the Royal Society B: Biological Sciences</i> , <b>2017</b> , 284,	4.4	20
63	Whole-exome sequencing of over 4100 men of African ancestry and prostate cancer risk. <i>Human Molecular Genetics</i> , <b>2016</b> , 25, 371-81	5.6	19
62	Assessing the performance of qpAdm: a statistical tool for studying population admixture. <i>Genetics</i> , <b>2021</b> , 217,	4	18

61	The Genomic History of the Bronze Age Southern Levant. <i>Cell</i> , <b>2020</b> , 181, 1146-1157.e11	56.2	17
60	A multi-stage genome-wide association study of uterine fibroids in African Americans. <i>Human Genetics</i> , <b>2017</b> , 136, 1363-1373	6.3	17
59	Human auditory ossicles as an alternative optimal source of ancient DNA. <i>Genome Research</i> , <b>2020</b> , 30, 427-436	9.7	16
58	Failure to replicate a genetic signal for sex bias in the steppe migration into central Europe. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2017</b> , 114, E3873-E3874	11.5	15
57	The Genomic Formation of South and Central Asia <b>2018</b> ,		15
56	The Genomic Formation of Human Populations in East Asia		14
55	Mitochondrial DNA analysis of eneolithic trypillians from Ukraine reveals neolithic farming genetic roots. <i>PLoS ONE</i> , <b>2017</b> , 12, e0172952	3.7	13
54	Interactions between earliest Linearbandkeramik farmers and central European hunter gatherers at the dawn of European Neolithization. <i>Scientific Reports</i> , <b>2019</b> , 9, 19544	4.9	13
53	Kinship and social organization in Copper Age Europe. A cross-disciplinary analysis of archaeology, DNA, isotopes, and anthropology from two Bell Beaker cemeteries. <i>PLoS ONE</i> , <b>2020</b> , 15, e0241278	3.7	12
52	Paleolithic DNA from the Caucasus reveals core of West Eurasian ancestry		12
51	Dairying enabled Early Bronze Age Yamnaya steppe expansions. <i>Nature</i> , <b>2021</b> , 598, 629-633	50.4	12
50	A Late Bronze Age II clay coffin from Tel Shaddud in the Central Jezreel Valley, Israel: context and historical implications. <i>Levant</i> , <b>2017</b> , 49, 105-135	0.5	11
49	Discussion: Are the Origins of Indo-European Languages Explained by the Migration of the Yamnaya Culture to the West?. <i>European Journal of Archaeology</i> , <b>2018</b> , 21, 3-17	0.7	10
48	Ancient DNA from the skeletons of Roopkund Lake reveals Mediterranean migrants in India. <i>Nature Communications</i> , <b>2019</b> , 10, 3670	17.4	10
47	Large-scale migration into Britain during the Middle to Late Bronze Age.. <i>Nature</i> , <b>2021</b> ,	50.4	10
46	No statistical evidence for an effect of CCR5-B2 on lifespan in the UK Biobank cohort. <i>Nature Medicine</i> , <b>2020</b> , 26, 178-180	50.5	10
45	Ethics of DNA research on human remains: five globally applicable guidelines. <i>Nature</i> , <b>2021</b> , 599, 41-46	50.4	9
44	ContamLD: estimation of ancient nuclear DNA contamination using breakdown of linkage disequilibrium. <i>Genome Biology</i> , <b>2020</b> , 21, 199	18.3	9



43	African Ancestry Analysis and Admixture Genetic Mapping for Proliferative Diabetic Retinopathy in African Americans <b>2015</b> , 56, 3999-4005		8
42	Ancient genomes in South Patagonia reveal population movements associated with technological shifts and geography. <i>Nature Communications</i> , <b>2020</b> , 11, 3868	17.4	8
41	Increased rate of close-kin unions in the central Andes in the half millennium before European contact. <i>Current Biology</i> , <b>2020</b> , 30, R980-R981	6.3	8
40	A high-resolution picture of kinship practices in an Early Neolithic tomb.. <i>Nature</i> , <b>2021</b> ,	50.4	7
39	Three Phases of Ancient Migration Shaped the Ancestry of Human Populations in Vanuatu. <i>Current Biology</i> , <b>2020</b> , 30, 4846-4856.e6	6.3	7
38	Ancient DNA reveals monozygotic newborn twins from the Upper Palaeolithic. <i>Communications Biology</i> , <b>2020</b> , 3, 650	6.7	6
37	The genomic formation of First American ancestors in East and Northeast Asia		6
36	No evidence for unknown archaic ancestry in South Asia. <i>Nature Genetics</i> , <b>2018</b> , 50, 632-633	36.3	5
35	The Kalash Genetic Isolate? The Evidence for Recent Admixture. <i>American Journal of Human Genetics</i> , <b>2016</b> , 98, 396-7	11	5
34	Population history from the Neolithic to present on the Mediterranean island of Sardinia: An ancient DNA perspective		5
33	The Arrival of Steppe and Iranian Related Ancestry in the Islands of the Western Mediterranean		5
32	A unified genealogy of modern and ancient genomes		4
31	South-to-north migration preceded the advent of intensive farming in the Maya region.. <i>Nature Communications</i> , <b>2022</b> , 13, 1530	17.4	4
30	Direct dating of human skeletal material from Ganj Dareh, Early Neolithic of the Iranian Zagros. <i>Journal of Archaeological Science: Reports</i> , <b>2017</b> , 12, 165-172	0.7	3
29	Genomic transformation and social organization during the Copper Age-Bronze Age transition in southern Iberia. <i>Science Advances</i> , <b>2021</b> , 7, eabi7038	14.3	3
28	ContamLD: Estimation of Ancient Nuclear DNA Contamination Using Breakdown of Linkage Disequilibrium		3
27	Assessing the Performance of qpAdm: A Statistical Tool for Studying Population Admixture		3
26	Two genetic variants explain the association of European ancestry with multiple sclerosis risk in African-Americans. <i>Scientific Reports</i> , <b>2020</b> , 10, 16902	4.9	3

25	A minimally destructive protocol for DNA extraction from ancient teeth. <i>Genome Research</i> , <b>2021</b> , 31, 472-483	9.7	3
24	Dynamic changes in genomic and social structures in third millennium BCE central Europe. <i>Science Advances</i> , <b>2021</b> , 7,	14.3	3
23	Stone Age genomes shed light on the early evolution, diversity, and ecology of plague.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2022</b> , 119, e2116722119	11.5	3
22	Revisiting ancient DNA insights into the human history of the Pacific Islands. <i>Archaeology in Oceania</i> , <b>2019</b> , 54, 53-56	0.7	2
21	Genetic landscape of Gullah African Americans. <i>American Journal of Physical Anthropology</i> , <b>2021</b> , 175, 905-919	2.5	2
20	Ancient mitochondrial genomes from the Argentinian Pampas inform the early peopling of the Southern Cone of South America. <i>iScience</i> , <b>2021</b> , 24, 102553	6.1	2
19	Social stratification without genetic differentiation at the site of Kulubnarti in Christian Period Nubia		2
18	The return of the Beaker folk? Rethinking migration and population change in British prehistory. <i>Antiquity</i> , 1-14	1	2
17	Mitochondrial genome diversity on the Central Siberian Plateau with particular reference to the prehistory of northernmost Eurasia. <i>PLoS ONE</i> , <b>2021</b> , 16, e0244228	3.7	2
16	Ancient DNA and deep population structure in sub-Saharan African foragers.. <i>Nature</i> , <b>2022</b> ,	50.4	2
15	Genetic Landscape of Gullah African Americans		1
14	Natural hybridization reveals incompatible alleles that cause melanoma in swordtail fish		1
13	Ancestry and demography and descendants of Iron Age nomads of the Eurasian Steppe		1
12	A genetic history of the pre-contact Caribbean		1
11	A Minimally Destructive Protocol for DNA Extraction from Ancient Teeth		1
10	Kinship and social organization in Copper Age Europe. A cross-disciplinary analysis of archaeology, DNA, isotopes, and anthropology from two Bell Beaker cemeteries		1
9	Genome-wide analysis of nearly all the victims of a 6200 year old massacre. <i>PLoS ONE</i> , <b>2021</b> , 16, e0247332	3.7	1
8	An ancient DNA Pacific journey: a case study of collaboration between archaeologists and geneticists. <i>World Archaeology</i> , <b>2019</b> , 51, 620-639	1.4	1

7	Cosmopolitanism at the Roman Danubian Frontier, Slavic Migrations, and the Genomic Formation of Modern Balkan Peoples		1
6	COMBINING ANCIENT DNA AND RADIOCARBON DATING DATA TO INCREASE CHRONOLOGICAL ACCURACY. <i>Journal of Archaeological Science</i> , <b>2021</b> , 133, 105452-105452	2.9	1
5	A unified genealogy of modern and ancient genomes.. <i>Science</i> , <b>2022</b> , 375, eabi8264	33.3	1
4	An integrative skeletal and paleogenomic analysis of stature variation suggests relatively reduced health for early European farmers.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2022</b> , 119, e2106743119	11.5	1
3	Social stratification without genetic differentiation at the site of Kulubnarti in Christian Period Nubia.. <i>Nature Communications</i> , <b>2021</b> , 12, 7283	17.4	0
2	Indian genetic heritage in Southeast Asian populations.. <i>PLoS Genetics</i> , <b>2022</b> , 18, e1010036	6	0
1	A High-Density Admixture Scan in 1,670 African Americans with Hypertension. <i>PLoS Genetics</i> , <b>2005</b> , preprint, e196	6	