

Nick Patterson

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

56
papers

22,913
citations

39
h-index

61
g-index

61
ext. papers

28,882
ext. citations

32.9
avg, IF

6.11
L-index

#	Paper	IF	Citations
56	Ancient DNA and deep population structure in sub-Saharan African foragers.. <i>Nature</i> , 2022 ,	50.4	2
55	A unified genealogy of modern and ancient genomes.. <i>Science</i> , 2022 , 375, eabi8264	33.3	1
54	Social stratification without genetic differentiation at the site of Kulubnarti in Christian Period Nubia.. <i>Nature Communications</i> , 2021 , 12, 7283	17.4	0
53	Large-scale migration into Britain during the Middle to Late Bronze Age.. <i>Nature</i> , 2021 ,	50.4	10
52	Ethics of DNA research on human remains: five globally applicable guidelines. <i>Nature</i> , 2021 , 599, 41-46	50.4	9
51	Investigate the origins of COVID-19. <i>Science</i> , 2021 , 372, 694	33.3	39
50	Assessing the performance of qpAdm: a statistical tool for studying population admixture. <i>Genetics</i> , 2021 , 217,	4	18
49	COMBINING ANCIENT DNA AND RADIOCARBON DATING DATA TO INCREASE CHRONOLOGICAL ACCURACY. <i>Journal of Archaeological Science</i> , 2021 , 133, 105452-105452	2.9	1
48	The spread of steppe and Iranian-related ancestry in the islands of the western Mediterranean. <i>Nature Ecology and Evolution</i> , 2020 , 4, 334-345	12.3	48
47	Ancient West African foragers in the context of African population history. <i>Nature</i> , 2020 , 577, 665-670	50.4	47
46	Two genetic variants explain the association of European ancestry with multiple sclerosis risk in African-Americans. <i>Scientific Reports</i> , 2020 , 10, 16902	4.9	3
45	ContamLD: estimation of ancient nuclear DNA contamination using breakdown of linkage disequilibrium. <i>Genome Biology</i> , 2020 , 21, 199	18.3	9
44	An Ancient Harappan Genome Lacks Ancestry from Steppe Pastoralists or Iranian Farmers. <i>Cell</i> , 2019 , 179, 729-735.e10	56.2	28
43	The formation of human populations in South and Central Asia. <i>Science</i> , 2019 , 365,	33.3	195
42	The genomic history of the Iberian Peninsula over the past 8000 years. <i>Science</i> , 2019 , 363, 1230-1234	33.3	186
41	Ancient DNA from the skeletons of Roopkund Lake reveals Mediterranean migrants in India. <i>Nature Communications</i> , 2019 , 10, 3670	17.4	10
40	Genomic architecture and introgression shape a butterfly radiation. <i>Science</i> , 2019 , 366, 594-599	33.3	161

39	The Beaker phenomenon and the genomic transformation of northwest Europe. <i>Nature</i> , 2018 , 555, 190-196	106	293
38	The genomic history of southeastern Europe. <i>Nature</i> , 2018 , 555, 197-203	50.4	287
37	Reconstructing the genetic history of late Neanderthals. <i>Nature</i> , 2018 , 555, 652-656	50.4	138
36	Ancient DNA from Chalcolithic Israel reveals the role of population mixture in cultural transformation. <i>Nature Communications</i> , 2018 , 9, 3336	17.4	37
35	The Genomic Formation of South and Central Asia 2018 ,		15
34	Reconstructing the Deep Population History of Central and South America. <i>Cell</i> , 2018 , 175, 1185-1197.e27	26.2	143
33	A high-coverage Neandertal genome from Vindija Cave in Croatia. <i>Science</i> , 2017 , 358, 655-658	33.3	312
32	Reconstructing Prehistoric African Population Structure. <i>Cell</i> , 2017 , 171, 59-71.e21	56.2	201
31	The promise of discovering population-specific disease-associated genes in South Asia. <i>Nature Genetics</i> , 2017 , 49, 1403-1407	36.3	79
30	Genetic origins of the Minoans and Mycenaeans. <i>Nature</i> , 2017 , 548, 214-218	50.4	108
29	The Simons Genome Diversity Project: 300 genomes from 142 diverse populations. <i>Nature</i> , 2016 , 538, 201-206	50.4	759
28	Genomic insights into the origin of farming in the ancient Near East. <i>Nature</i> , 2016 , 536, 419-24	50.4	485
27	The Kalash Genetic Isolate? The Evidence for Recent Admixture. <i>American Journal of Human Genetics</i> , 2016 , 98, 396-7	11	5
26	The genetic history of Ice Age Europe. <i>Nature</i> , 2016 , 534, 200-5	50.4	473
25	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> , 2016 , 113, 5652-7	11.5	75
24	Genomic insights into the peopling of the Southwest Pacific. <i>Nature</i> , 2016 , 538, 510-513	50.4	180
23	Massive migration from the steppe was a source for Indo-European languages in Europe. <i>Nature</i> , 2015 , 522, 207-11	50.4	968
22	Genetic evidence for two founding populations of the Americas. <i>Nature</i> , 2015 , 525, 104-8	50.4	220

21	An early modern human from Romania with a recent Neanderthal ancestor. <i>Nature</i> , 2015 , 524, 216-9	50.4	446
20	Genome-wide patterns of selection in 230 ancient Eurasians. <i>Nature</i> , 2015 , 528, 499-503	50.4	774
19	The complete genome sequence of a Neanderthal from the Altai Mountains. <i>Nature</i> , 2014 , 505, 43-9	50.4	1339
18	The genomic landscape of Neanderthal ancestry in present-day humans. <i>Nature</i> , 2014 , 507, 354-7	50.4	615
17	Ancient human genomes suggest three ancestral populations for present-day Europeans. <i>Nature</i> , 2014 , 513, 409-13	50.4	812
16	Genetic evidence for recent population mixture in India. <i>American Journal of Human Genetics</i> , 2013 , 93, 422-38	11	177
15	Inferring admixture histories of human populations using linkage disequilibrium. <i>Genetics</i> , 2013 , 193, 1233-54	4	293
14	Ancient admixture in human history. <i>Genetics</i> , 2012 , 192, 1065-93	4	1212
13	A high-coverage genome sequence from an archaic Denisovan individual. <i>Science</i> , 2012 , 338, 222-6	33.3	1276
12	Reconstructing Native American population history. <i>Nature</i> , 2012 , 488, 370-4	50.4	498
11	The genetic prehistory of southern Africa. <i>Nature Communications</i> , 2012 , 3, 1143	17.4	193
10	Denisova admixture and the first modern human dispersals into Southeast Asia and Oceania. <i>American Journal of Human Genetics</i> , 2011 , 89, 516-28	11	390
9	The landscape of recombination in African Americans. <i>Nature</i> , 2011 , 476, 170-5	50.4	243
8	The history of African gene flow into Southern Europeans, Levantines, and Jews. <i>PLoS Genetics</i> , 2011 , 7, e1001373	6	175
7	Genetic history of an archaic hominin group from Denisova Cave in Siberia. <i>Nature</i> , 2010 , 468, 1053-60	50.4	1169
6	Reconstructing Indian population history. <i>Nature</i> , 2009 , 461, 489-94	50.4	1075
5	A second generation human haplotype map of over 3.1 million SNPs. <i>Nature</i> , 2007 , 449, 851-61	50.4	3647
4	Population structure and eigenanalysis. <i>PLoS Genetics</i> , 2006 , 2, e190	6	3021

3	Genomic architecture and introgression shape a butterfly radiation	7
2	Social stratification without genetic differentiation at the site of Kulubnarti in Christian Period Nubia	2
1	A unified genealogy of modern and ancient genomes	4