

Eadaoin Harney

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2100763/publications.pdf>

Version: 2024-02-01

23
papers

6,883
citations

361045

20
h-index

610482

24
g-index

38
all docs

38
docs citations

38
times ranked

5719
citing authors

#	ARTICLE	IF	CITATIONS
1	Massive migration from the steppe was a source for Indo-European languages in Europe. <i>Nature</i> , 2015, 522, 207-211.	13.7	1,435
2	Genome-wide patterns of selection in 230 ancient Eurasians. <i>Nature</i> , 2015, 528, 499-503.	13.7	1,160
3	Genomic insights into the origin of farming in the ancient Near East. <i>Nature</i> , 2016, 536, 419-424.	13.7	733
4	The Beaker phenomenon and the genomic transformation of northwest Europe. <i>Nature</i> , 2018, 555, 190-196.	13.7	503
5	The genomic history of southeastern Europe. <i>Nature</i> , 2018, 555, 197-203.	13.7	479
6	The formation of human populations in South and Central Asia. <i>Science</i> , 2019, 365, .	6.0	383
7	Partial uracilâ€“DNAâ€“glycosylase treatment for screening of ancient DNA. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2015, 370, 20130624.	1.8	381
8	Reconstructing Prehistoric African Population Structure. <i>Cell</i> , 2017, 171, 59-71.e21.	13.5	308
9	Parallel palaeogenomic transects reveal complex genetic history of early European farmers. <i>Nature</i> , 2017, 551, 368-372.	13.7	306
10	Genomic insights into the peopling of the Southwest Pacific. <i>Nature</i> , 2016, 538, 510-513.	13.7	262
11	Ancient genomes document multiple waves of migration in Southeast Asian prehistory. <i>Science</i> , 2018, 361, 92-95.	6.0	250
12	Ancient human genome-wide data from a 3000-year interval in the Caucasus corresponds with eco-geographic regions. <i>Nature Communications</i> , 2019, 10, 590.	5.8	113
13	Assessing the performance of qpAdm: a statistical tool for studying population admixture. <i>Genetics</i> , 2021, 217, .	1.2	84
14	Ancient DNA from Chalcolithic Israel reveals the role of population mixture in cultural transformation. <i>Nature Communications</i> , 2018, 9, 3336.	5.8	71
15	A Paleogenomic Reconstruction of the Deep Population History of the Andes. <i>Cell</i> , 2020, 181, 1131-1145.e21.	13.5	69
16	Multi-layered population structure in Island Southeast Asians. <i>European Journal of Human Genetics</i> , 2016, 24, 1605-1611.	1.4	50
17	Human auditory ossicles as an alternative optimal source of ancient DNA. <i>Genome Research</i> , 2020, 30, 427-436.	2.4	37
18	A minimally destructive protocol for DNA extraction from ancient teeth. <i>Genome Research</i> , 2021, 31, 472-483.	2.4	31

#	ARTICLE	IF	CITATIONS
19	Stone Age <i>Yersinia pestis</i> genomes shed light on the early evolution, diversity, and ecology of plague. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, e2116722119.	3.3	31
20	ContamLD: estimation of ancient nuclear DNA contamination using breakdown of linkage disequilibrium. Genome Biology, 2020, 21, 199.	3.8	29
21	Ancient DNA from the skeletons of Roopkund Lake reveals Mediterranean migrants in India. Nature Communications, 2019, 10, 3670.	5.8	19
22	Social stratification without genetic differentiation at the site of Kulubnarti in Christian Period Nubia. Nature Communications, 2021, 12, 7283.	5.8	13
23	Combining ancient DNA and radiocarbon dating data to increase chronological accuracy. Journal of Archaeological Science, 2021, 133, 105452.	1.2	7