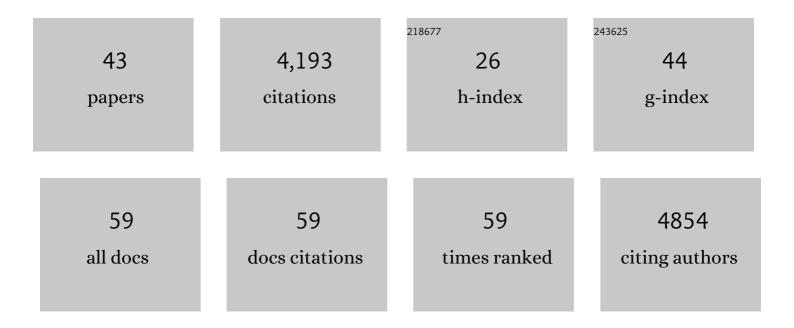
Iñigo Olalde

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3669600/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	A high-resolution picture of kinship practices in an Early Neolithic tomb. Nature, 2022, 601, 584-587.	27.8	65
2	Large-scale migration into Britain during the Middle to Late Bronze Age. Nature, 2022, 601, 588-594.	27.8	86
3	Ancient DNA and deep population structure in sub-Saharan African foragers. Nature, 2022, 603, 290-296.	27.8	51
4	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.	7.1	31
5	At the beginnings of the funerary Megalithism in Iberia at Campo de Hockey necropolis. Scientific Reports, 2022, 12, .	3.3	8
6	A genetic history of the pre-contact Caribbean. Nature, 2021, 590, 103-110.	27.8	67
7	Genome-wide analysis of nearly all the victims of a 6200 year old massacre. PLoS ONE, 2021, 16, e0247332.	2.5	11
8	Combining ancient DNA and radiocarbon dating data to increase chronological accuracy. Journal of Archaeological Science, 2021, 133, 105452.	2.4	7
9	Salmonella enterica from a soldier from the 1652 siege of Barcelona (Spain) supports historical transatlantic epidemic contacts. IScience, 2021, 24, 103021.	4.1	2
10	Genomic transformation and social organization during the Copper Age–Bronze Age transition in southern Iberia. Science Advances, 2021, 7, eabi7038.	10.3	39
11	Social stratification without genetic differentiation at the site of Kulubnarti in Christian Period Nubia. Nature Communications, 2021, 12, 7283.	12.8	13
12	Genomic Analysis of 18th-Century Kazakh Individuals and Their Oral Microbiome. Biology, 2021, 10, 1324.	2.8	2
13	Plasmodium vivax Malaria Viewed through the Lens of an Eradicated European Strain. Molecular Biology and Evolution, 2020, 37, 773-785.	8.9	38
14	Ancient DNA reveals monozygotic newborn twins from the Upper Palaeolithic. Communications Biology, 2020, 3, 650.	4.4	25
15	Latest trends in archaeogenetic research of west Eurasians. Current Opinion in Genetics and Development, 2020, 62, 36-43.	3.3	16
16	The spread of steppe and Iranian-related ancestry in the islands of the western Mediterranean. Nature Ecology and Evolution, 2020, 4, 334-345.	7.8	95
17	Ancient West African foragers in the context of African population history. Nature, 2020, 577, 665-670.	27.8	86
18	Kinship and social organization in Copper Age Europe. A cross-disciplinary analysis of archaeology, DNA, isotopes, and anthropology from two Bell Beaker cemeteries. PLoS ONE, 2020, 15, e0241278.	2.5	35

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19	The formation of human populations in South and Central Asia. Science, 2019, 365, .	12.6	383
20	Evolutionary and functional impact of common polymorphic inversions in the human genome. Nature Communications, 2019, 10, 4222.	12.8	34
21	Ancient DNA reveals a multistep spread of the first herders into sub-Saharan Africa. Science, 2019, 365,	12.6	96
22	Palaeo-Eskimo genetic ancestry and the peopling of Chukotka and North America. Nature, 2019, 570, 236-240.	27.8	118
23	Ancient genomes indicate population replacement in Early Neolithic Britain. Nature Ecology and Evolution, 2019, 3, 765-771.	7.8	156
24	The genomic history of the Iberian Peninsula over the past 8000 years. Science, 2019, 363, 1230-1234.	12.6	340
25	The Beaker phenomenon and the genomic transformation of northwest Europe. Nature, 2018, 555, 190-196.	27.8	503
26	The genomic history of southeastern Europe. Nature, 2018, 555, 197-203.	27.8	479
27	On the path to extinction: Inbreeding and admixture in a declining grey wolf population. Molecular Ecology, 2018, 27, 3599-3612.	3.9	46
28	Malaria was a weak selective force in ancient Europeans. Scientific Reports, 2017, 7, 1377.	3.3	32
29	Reconstructing Prehistoric African Population Structure. Cell, 2017, 171, 59-71.e21.	28.9	308
30	Mitochondrial DNA from the eradicated European <i>Plasmodium vivax</i> and <i>P. falciparum</i> from 70-year-old slides from the Ebro Delta in Spain. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 11495-11500.	7.1	41
31	Ancient DNA sheds light on the ancestry of pre-hispanic Canarian pigs. Genetics Selection Evolution, 2015, 47, 40.	3.0	13
32	Modern humans' paleogenomics and the new evidences on the European prehistory. Science and Technology of Archaeological Research, 2015, 1, 1-9.	2.4	5
33	Genome data from a sixteenth century pig illuminate modern breed relationships. Heredity, 2015, 114, 175-184.	2.6	39
34	A Common Genetic Origin for Early Farmers from Mediterranean Cardial and Central European LBK Cultures. Molecular Biology and Evolution, 2015, 32, msv181.	8.9	155
35	Limiting replication stress during somatic cell reprogramming reduces genomic instability in induced pluripotent stem cells. Nature Communications, 2015, 6, 8036.	12.8	84
36	Genomic analysis of the blood attributed to Louis XVI (1754–1793), king of France. Scientific Reports, 2015, 4, 4666.	3.3	16

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37	Mitochondrial DNA from El Mirador Cave (Atapuerca, Spain) Reveals the Heterogeneity of Chalcolithic Populations. PLoS ONE, 2014, 9, e105105.	2.5	28
38	Derived immune and ancestral pigmentation alleles in a 7,000-year-old Mesolithic European. Nature, 2014, 507, 225-228.	27.8	328
39	Analysis of structural diversity in wolf-like canids reveals post-domestication variants. BMC Genomics, 2014, 15, 465.	2.8	16
40	Population connectivity buffers genetic diversity loss in a seabird. Frontiers in Zoology, 2013, 10, 28.	2.0	20
41	Genetic comparison of the head of Henri IV and the presumptive blood from Louis XVI (both Kings of) Tj ETQq1 1	0,784314	l rgBT /Overl
42	Genomic Affinities of Two 7,000-Year-Old Iberian Hunter-Gatherers. Current Biology, 2012, 22, 1494-1499.	3.9	160
43	â€~Ava': a Beaker-associated woman from a cist at Achavanich, Highland, and the story of her (re-)discovery and subsequent study. Proceedings of the Society of Antiquaries of Scotland, 0, 147, 73-118.	0.0	3