

Iñigo Olalde

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3669600/publications.pdf>

Version: 2024-02-01

43
papers

4,193
citations

218677

26
h-index

243625

44
g-index

59
all docs

59
docs citations

59
times ranked

4854
citing authors

#	ARTICLE	IF	CITATIONS
19	The formation of human populations in South and Central Asia. <i>Science</i> , 2019, 365, .	12.6	383
20	Evolutionary and functional impact of common polymorphic inversions in the human genome. <i>Nature Communications</i> , 2019, 10, 4222.	12.8	34
21	Ancient DNA reveals a multistep spread of the first herders into sub-Saharan Africa. <i>Science</i> , 2019, 365, .	12.6	96
22	Palaeo-Eskimo genetic ancestry and the peopling of Chukotka and North America. <i>Nature</i> , 2019, 570, 236-240.	27.8	118
23	Ancient genomes indicate population replacement in Early Neolithic Britain. <i>Nature Ecology and Evolution</i> , 2019, 3, 765-771.	7.8	156
24	The genomic history of the Iberian Peninsula over the past 8000 years. <i>Science</i> , 2019, 363, 1230-1234.	12.6	340
25	The Beaker phenomenon and the genomic transformation of northwest Europe. <i>Nature</i> , 2018, 555, 190-196.	27.8	503
26	The genomic history of southeastern Europe. <i>Nature</i> , 2018, 555, 197-203.	27.8	479
27	On the path to extinction: Inbreeding and admixture in a declining grey wolf population. <i>Molecular Ecology</i> , 2018, 27, 3599-3612.	3.9	46
28	Malaria was a weak selective force in ancient Europeans. <i>Scientific Reports</i> , 2017, 7, 1377.	3.3	32
29	Reconstructing Prehistoric African Population Structure. <i>Cell</i> , 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. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 11495-11500.	7.1	41
31	Ancient DNA sheds light on the ancestry of pre-hispanic Canarian pigs. <i>Genetics Selection Evolution</i> , 2015, 47, 40.	3.0	13
32	Modern humans' paleogenomics and the new evidences on the European prehistory. <i>Science and Technology of Archaeological Research</i> , 2015, 1, 1-9.	2.4	5
33	Genome data from a sixteenth century pig illuminate modern breed relationships. <i>Heredity</i> , 2015, 114, 175-184.	2.6	39
34	A Common Genetic Origin for Early Farmers from Mediterranean Cardial and Central European LBK Cultures. <i>Molecular Biology and Evolution</i> , 2015, 32, msv181.	8.9	155
35	Limiting replication stress during somatic cell reprogramming reduces genomic instability in induced pluripotent stem cells. <i>Nature Communications</i> , 2015, 6, 8036.	12.8	84
36	Genomic analysis of the blood attributed to Louis XVI (1754-1793), king of France. <i>Scientific Reports</i> , 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 France) reveals a common ancestor. PLoS ONE, 2014, 9, e105105.	2.2	13
42	Genomic Affinities of Two 7,000-Year-Old Iberian Hunter-Gatherers. Current Biology, 2012, 22, 1494-1499.	3.9	160
43	“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, 2014, 147, 73-118.	0.0	3