

François Xavier Ricaut

List of Publications by Year in descending order

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Version: 2024-02-01

47
papers

3,618
citations

236925

25
h-index

223800

46
g-index

48
all docs

48
docs citations

48
times ranked

5095
citing authors

#	ARTICLE	IF	CITATIONS
1	An Aboriginal Australian Genome Reveals Separate Human Dispersals into Asia. <i>Science</i> , 2011, 334, 94-98.	12.6	675
2	A genomic history of Aboriginal Australia. <i>Nature</i> , 2016, 538, 207-214.	27.8	439
3	Genomic analyses inform on migration events during the peopling of Eurasia. <i>Nature</i> , 2016, 538, 238-242.	27.8	360
4	A recent bottleneck of Y chromosome diversity coincides with a global change in culture. <i>Genome Research</i> , 2015, 25, 459-466.	5.5	348
5	Pig Domestication and Human-Mediated Dispersal in Western Eurasia Revealed through Ancient DNA and Geometric Morphometrics. <i>Molecular Biology and Evolution</i> , 2013, 30, 824-832.	8.9	196
6	Multiple Deeply Divergent Denisovan Ancestries in Papuans. <i>Cell</i> , 2019, 177, 1010-1021.e32.	28.9	181
7	Ancient DNA reveals male diffusion through the Neolithic Mediterranean route. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 9788-9791.	7.1	151
8	Ancient DNA suggests the leading role played by men in the Neolithic dissemination. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 18255-18259.	7.1	103
9	Ancient pigs reveal a near-complete genomic turnover following their introduction to Europe. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 17231-17238.	7.1	101
10	Genomic landscape of human diversity across Madagascar. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, E6498-E6506.	7.1	77
11	Genome-wide evidence of Austronesian-Bantu admixture and cultural reversion in a hunter-gatherer group of Madagascar. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 936-941.	7.1	75
12	Selective sweep on human amylase genes postdates the split with Neanderthals. <i>Scientific Reports</i> , 2016, 6, 37198.	3.3	67
13	Malagasy Genetic Ancestry Comes from an Historical Malay Trading Post in Southeast Borneo. <i>Molecular Biology and Evolution</i> , 2016, 33, 2396-2400.	8.9	62
14	Mitochondrial DNA and the Y chromosome suggest the settlement of Madagascar by Indonesian sea nomad populations. <i>BMC Genomics</i> , 2015, 16, 191.	2.8	61
15	STR-genotyping from human medieval tooth and bone samples. <i>Forensic Science International</i> , 2005, 151, 31-35.	2.2	58
16	Strong selection during the last millennium for African ancestry in the admixed population of Madagascar. <i>Nature Communications</i> , 2018, 9, 932.	12.8	57
17	Population structure of modern-day Italians reveals patterns of ancient and archaic ancestries in Southern Europe. <i>Science Advances</i> , 2019, 5, eaaw3492.	10.3	53
18	Multi-layered population structure in Island Southeast Asians. <i>European Journal of Human Genetics</i> , 2016, 24, 1605-1611.	2.8	50

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19	Contrasting Linguistic and Genetic Origins of the Asian Source Populations of Malagasy. <i>Scientific Reports</i> , 2016, 6, 26066.	3.3	48
20	Genome-Wide Characterization of Arabian Peninsula Populations: Shedding Light on the History of a Fundamental Bridge between Continents. <i>Molecular Biology and Evolution</i> , 2019, 36, 575-586.	8.9	45
21	Genetic analysis and ethnic affinities from two Scytho-Siberian skeletons. <i>American Journal of Physical Anthropology</i> , 2004, 123, 351-360.	2.1	43
22	Tracing Arab-Islamic Inheritance in Madagascar: Study of the Y-chromosome and Mitochondrial DNA in the Antemoro. <i>PLoS ONE</i> , 2013, 8, e80932.	2.5	42
23	The Comoros Show the Earliest Austronesian Gene Flow into the Swahili Corridor. <i>American Journal of Human Genetics</i> , 2018, 102, 58-68.	6.2	32
24	Ancient DNA analysis of human neolithic remains found in northeastern Siberia. <i>American Journal of Physical Anthropology</i> , 2005, 126, 458-462.	2.1	29
25	Molecular Genetic Analysis of 400-Year-Old Human Remains Found in Two Yakut Burial Sites. <i>American Journal of Physical Anthropology</i> , 2006, 129, 55-63.	2.1	25
26	Papuan mitochondrial genomes and the settlement of Sahul. <i>Journal of Human Genetics</i> , 2020, 65, 875-887.	2.3	24
27	The last sea nomads of the Indonesian archipelago: genomic origins and dispersal. <i>European Journal of Human Genetics</i> , 2017, 25, 1004-1010.	2.8	21
28	Molecular Identification of Bacteria by Total Sequence Screening: Determining the Cause of Death in Ancient Human Subjects. <i>PLoS ONE</i> , 2011, 6, e21733.	2.5	19
29	Genetic analysis of human remains found in two eighteenth century Yakut graves at At-Dabaan. <i>International Journal of Legal Medicine</i> , 2004, 118, 24-31.	2.2	18
30	Evidence of Austronesian Genetic Lineages in East Africa and South Arabia: Complex Dispersal from Madagascar and Southeast Asia. <i>Genome Biology and Evolution</i> , 2019, 11, 748-758.	2.5	15
31	Cranial Discrete Traits in a Byzantine Population and Eastern Mediterranean Population Movements. <i>Human Biology</i> , 2008, 80, 535-564.	0.2	14
32	New insights on the late Pleistocene–Holocene lithic industry in East Kalimantan (Borneo): The contribution of three rock shelter sites in the karstic area of the Mangkalihat peninsula. <i>Quaternary International</i> , 2016, 416, 126-150.	1.5	14
33	Testing for kavain in human hair using gas chromatography–tandem mass spectrometry. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2003, 798, 351-354.	2.3	13
34	Genomic admixture tracks pulses of economic activity over 2,000 years in the Indian Ocean trading network. <i>Scientific Reports</i> , 2017, 7, 2919.	3.3	13
35	A Time Series of Prehistoric Mitochondrial DNA Reveals Western European Genetic Diversity Was Largely Established by the Bronze Age. <i>Advances in Anthropology</i> , 2012, 02, 14-23.	0.2	12
36	Papua New Guinean Genomes Reveal the Complex Settlement of North Sahul. <i>Molecular Biology and Evolution</i> , 2021, 38, 5107-5121.	8.9	11

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37	Ancient Solomon Islands mtDNA: assessing Holocene settlement and the impact of European contact. <i>Journal of Archaeological Science</i> , 2010, 37, 1161-1170.	2.4	10
38	Episodes of Diversification and Isolation in Island Southeast Asian and Near Oceanian Male Lineages. <i>Molecular Biology and Evolution</i> , 2022, 39, .	8.9	9
39	Mitochondrial DNA Variation in Karkar Islanders. <i>Annals of Human Genetics</i> , 2008, 72, 349-367.	0.8	8
40	Western Eurasian genetic influences in the Indonesian archipelago. <i>Quaternary International</i> , 2016, 416, 243-248.	1.5	8
41	Tissue- and ethnicity-independent hypervariable DNA methylation states show evidence of establishment in the early human embryo. <i>Nucleic Acids Research</i> , 2022, 50, 6735-6752.	14.5	8
42	Testing for Betel Nut Alkaloids in Hair of Papuans Abusers using UPLC-MS/MS and UPLC-Q-ToF-MS. <i>Journal of Analytical Toxicology</i> , 2020, 44, 41-48.	2.8	6
43	Borneo as a half empty pot: Pottery assemblage from Liang Abu, East Kalimantan. <i>Quaternary International</i> , 2016, 416, 228-242.	1.5	4
44	Phenotypic differences between highlanders and lowlanders in Papua New Guinea. <i>PLoS ONE</i> , 2021, 16, e0253921.	2.5	4
45	Rock Art and (Re)Production of Narratives: A Cassowary Bone Dagger Stencil Perspective from Auwim, East Sepik, Papua New Guinea. <i>Cambridge Archaeological Journal</i> , 2022, 32, 547-565.	0.9	3
46	Chronology of natural selection in Oceanian genomes. <i>IScience</i> , 2022, 25, 104583.	4.1	3
47	mtDNA variation in the Buryat population of the Barguzin Valley: New insights into the micro-evolutionary history of the Baikal area. <i>Annals of Human Biology</i> , 2010, 37, 501-523.	1.0	2