FranÃ§ois Xavier Ricaut

List of Publications by Year
in descending order

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	Rock Art and (Re)Production of Narratives: A Cassowary Bone Dagger Stencil Perspective from Auwim, East Sepik, Papua New Guinea. Cambridge Archaeological Journal, 2022, 32, 547-565.	0.9	3
2	Episodes of Diversification and Isolation in Island Southeast Asian and Near Oceanian Male Lineages. Molecular Biology and Evolution, 2022, 39, .	8.9	9
3	Tissue- and ethnicity-independent hypervariable DNA methylation states show evidence of establishment in the early human embryo. Nucleic Acids Research, 2022, 50, 6735-6752.	14.5	8
4	Chronology of natural selection in Oceanian genomes. IScience, 2022, 25, 104583.	4.1	3
5	Phenotypic differences between highlanders and lowlanders in Papua New Guinea. PLoS ONE, 2021, 16, e0253921.	2.5	4
6	Papua New Guinean Genomes Reveal the Complex Settlement of North Sahul. Molecular Biology and Evolution, 2021, 38, 5107-5121.	8.9	11
7	Testing for Betel Nut Alkaloids in Hair of Papuans Abusers using UPLCâ€“MS/MS and UPLCâ€“Q-Tof-MS. Journal of Analytical Toxicology, 2020, 44, 41-48.	2.8	6
8	Papuan mitochondrial genomes and the settlement of Sahul. Journal of Human Genetics, 2020, 65, 875-887.	2.3	24
9	Ancient pigs reveal a near-complete genomic turnover following their introduction to Europe. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 17231-17238.	7.1	101
10	Population structure of modern-day Italians reveals patterns of ancient and archaic ancestries in Southern Europe. Science Advances, 2019, 5, eaaw3492.	10.3	53
11	Multiple Deeply Divergent Denisovan Ancestries in Papuans. Cell, 2019, 177, 1010-1021.e32.	28.9	181
12	Genome-Wide Characterization of Arabian Peninsula Populations: Shedding Light on the History of a Fundamental Bridge between Continents. Molecular Biology and Evolution, 2019, 36, 575-586.	8.9	45
13	Evidence of Austronesian Genetic Lineages in East Africa and South Arabia: Complex Dispersal from Madagascar and Southeast Asia. Genome Biology and Evolution, 2019, 11, 748-758.	2.5	15
14	The Comoros Show the Earliest Austronesian Gene Flow into the Swahili Corridor. American Journal of Human Genetics, 2018, 102, 58-68.	6.2	32
15	Strong selection during the last millennium for African ancestry in the admixed population of Madagascar. Nature Communications, 2018, 9, 932.	12.8	57
16	The last sea nomads of the Indonesian archipelago: genomic origins and dispersal. European Journal of Human Genetics, 2017, 25, 1004-1010.	2.8	21
17	Genomic landscape of human diversity across Madagascar. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E6498-E6506.	7.1	77
18	Genomic admixture tracks pulses of economic activity over 2,000 years in the Indian Ocean trading network. Scientific Reports, 2017, 7, 2919.	3.3	13

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19	Malagasy Genetic Ancestry Comes from an Historical Malay Trading Post in Southeast Borneo. Molecular Biology and Evolution, 2016, 33, 2396-2400.	8.9	62
20	New insights on the late Pleistoceneâ€“Holocene lithic industry in East Kalimantan (Borneo): The contribution of three rock shelter sites in the karstic area of the Mangkalihat peninsula. Quaternary International, 2016, 416, 126-150.	1.5	14
21	A genomic history of Aboriginal Australia. Nature, 2016, 538, 207-214.	27.8	439
22	Genomic analyses inform on migration events during the peopling of Eurasia. Nature, 2016, 538, 238-242.	27.8	360
23	Contrasting Linguistic and Genetic Origins of the Asian Source Populations of Malagasy. Scientific Reports, 2016, 6, 26066.	3.3	48
24	Selective sweep on human amylase genes postdates the split with Neanderthals. Scientific Reports, 2016, 6, 37198.	3.3	67
25	Multi-layered population structure in Island Southeast Asians. European Journal of Human Genetics, 2016, 24, 1605-1611.	2.8	50
26	Borneo as a half empty pot: Pottery assemblage from Liang Abu, East Kalimantan. Quaternary International, 2016, 416, 228-242.	1.5	4
27	Western Eurasian genetic influences in the Indonesian archipelago. Quaternary International, 2016, 416, 243-248.	1.5	8
28	Mitochondrial DNA and the Y chromosome suggest the settlement of Madagascar by Indonesian sea nomad populations. BMC Genomics, 2015, 16, 191.	2.8	61
29	A recent bottleneck of Y chromosome diversity coincides with a global change in culture. Genome Research, 2015, 25, 459-466.	5.5	348
30	Genome-wide evidence of Austronesianâ€“Bantu admixture and cultural reversion in a hunter-gatherer group of Madagascar. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 936-941.	7.1	75
31	Pig Domestication and Human-Mediated Dispersal in Western Eurasia Revealed through Ancient DNA and Geometric Morphometrics. Molecular Biology and Evolution, 2013, 30, 824-832.	8.9	196
32	Tracing Arab-Islamic Inheritance in Madagascar: Study of the Y-chromosome and Mitochondrial DNA in the Antemoro. PLoS ONE, 2013, 8, e80932.	2.5	42
33	A Time Series of Prehistoric Mitochondrial DNA Reveals Western European Genetic Diversity Was Largely Established by the Bronze Age. Advances in Anthropology, 2012, 02, 14-23.	0.2	12
34	Molecular Identification of Bacteria by Total Sequence Screening: Determining the Cause of Death in Ancient Human Subjects. PLoS ONE, 2011, 6, e21733.	2.5	19
35	Ancient DNA reveals male diffusion through the Neolithic Mediterranean route. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 9788-9791.	7.1	151
36	Ancient DNA suggests the leading role played by men in the Neolithic dissemination. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 18255-18259.	7.1	103

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37	An Aboriginal Australian Genome Reveals Separate Human Dispersals into Asia. Science, 2011, 334, 94-98.	12.6	675
38	Ancient Solomon Islands mtDNA: assessing Holocene settlement and the impact of European contact. Journal of Archaeological Science, 2010, 37, 1161-1170.	2.4	10
39	mtDNA variation in the Buryat population of the Barguzin Valley: New insights into the micro-evolutionary history of the Baikal area. Annals of Human Biology, 2010, 37, 501-523.	1.0	2
40	Mitochondrial DNA Variation in Karkar Islanders. Annals of Human Genetics, 2008, 72, 349-367.	0.8	8
41	Cranial Discrete Traits in a Byzantine Population and Eastern Mediterranean Population Movements. Human Biology, 2008, 80, 535-564.	0.2	14
42	Molecular Genetic Analysis of 400-Year-Old Human Remains Found in Two Yakut Burial Sites. American Journal of Physical Anthropology, 2006, 129, 55-63.	2.1	25
43	Ancient DNA analysis of human neolithic remains found in northeastern Siberia. American Journal of Physical Anthropology, 2005, 126, 458-462.	2.1	29
44	STR-genotyping from human medieval tooth and bone samples. Forensic Science International, 2005, 151, 31-35.	2.2	58
45	Genetic analysis of human remains found in two eighteenth century Yakut graves at At-Dabaan. International Journal of Legal Medicine, 2004, 118, 24-31.	2.2	18
46	Genetic analysis and ethnic affinities from two Scytho-Siberian skeletons. American Journal of Physical Anthropology, 2004, 123, 351-360.	2.1	43
47	Testing for kavain in human hair using gas chromatographyâ€“tandem mass spectrometry. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2003, 798, 351-354.	2.3	13