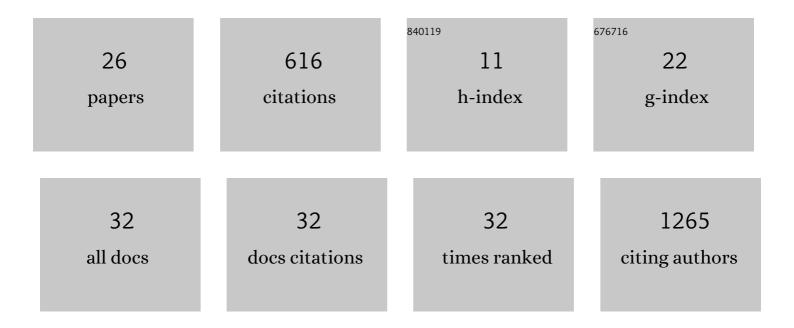
Cesar Fortes-Lima

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6850440/publications.pdf

Version: 2024-02-01



#	Article	IF	CITATIONS
1	Closing the Gaps in Genomic Research. Trends in Genetics, 2021, 37, 104-106.	2.9	4
2	Demographic history and admixture dynamics in African Sahelian populations. Human Molecular Genetics, 2021, 30, R29-R36.	1.4	8
3	Complex genetic admixture histories reconstructed with Approximate Bayesian Computation. Molecular Ecology Resources, 2021, 21, 1098-1117.	2.2	13
4	Population collapse in Congo rainforest from 400 CE urges reassessment of the Bantu Expansion. Science Advances, 2021, 7, .	4.7	30
5	Genetic substructure and complex demographic history of South African Bantu speakers. Nature Communications, 2021, 12, 2080.	5.8	47
6	Identification of ancestry proportions in admixed groups across the Americas using clinical pharmacogenomic SNP panels. Scientific Reports, 2021, 11, 1007.	1.6	5
7	Anthropological genetics perspectives on the transatlantic slave trade. Human Molecular Genetics, 2021, 30, R79-R87.	1.4	18
8	Engagement Cross-Disciplinary Research in Africa. , 2021, , 1-6.		0
9	Disentangling the Impact of the Transatlantic Slave Trade in African Diaspora Populations from a Genomic Perspective. , 2021, , 305-328.		0
10	Along the Indian Ocean Coast: Genomic Variation in Mozambique Provides New Insights into the Bantu Expansion. Molecular Biology and Evolution, 2020, 37, 406-416.	3.5	32
11	Predicting haplogroups using a versatile machine learning program (PredYMaLe) on a new mutationally balanced 32 Y-STR multiplex (CombYplex): Unlocking the full potential of the human STR mutation rate spectrum to estimate forensic parameters. Forensic Science International: Genetics, 2020, 48, 102342.	1.6	7
12	Novel insights on demographic history of tribal and caste groups from West Maharashtra (India) using genome-wide data. Scientific Reports, 2020, 10, 10075.	1.6	9
13	Meta-analysis of GWA studies provides new insights on the genetic architecture of skin pigmentation in recently admixed populations. BMC Genetics, 2019, 20, 59.	2.7	32
14	Unraveling African diversity from a crossâ€disciplinary perspective. Evolutionary Anthropology, 2019, 28, 288-292.	1.7	1
15	Reconstructing an African haploid genome from the 18th century. Nature Genetics, 2018, 50, 199-205.	9.4	15
16	Exploring Cuba's population structure and demographic history using genome-wide data. Scientific Reports, 2018, 8, 11422.	1.6	31
17	Dispersals and genetic adaptation of Bantu-speaking populations in Africa and North America. Science, 2017, 356, 543-546.	6.0	188
18	Genome-wide Ancestry and Demographic History of African-Descendant Maroon Communities from French Guiana and Suriname. American Journal of Human Genetics, 2017, 101, 725-736.	2.6	50

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#	Article	IF	CITATIONS
19	Revisiting Genetic Ancestry in African Diaspora Communities from Atlantic South America. , 2017, , 9-17.		0
20	Genetic population study of Y-chromosome markers in Benin and Ivory Coast ethnic groups. Forensic Science International: Genetics, 2015, 19, 232-237.	1.6	7
21	Impact of climate change and man-made irrigation systems on the transmission risk, long-term trend and seasonality of human and animal fascioliasis in Pakistan. Geospatial Health, 2014, 8, 317.	0.3	76
22	New insights into the distribution of <i>APOE</i> polymorphism in the Iberian Peninsula. The case of Andalusia (Spain). Annals of Human Biology, 2014, 41, 443-452.	0.4	5
23	Immunoglobulin genes in Andalusia (Spain). Genetic diversity in the Mediterranean space. Comptes Rendus - Biologies, 2014, 337, 646-656.	0.1	9
24	Y-STR genetic diversity in autochthonous Andalusians from Huelva and Granada provinces (Spain). Forensic Science International: Genetics, 2012, 6, e66-e71.	1.6	9
25	The Andalusian population from Huelva reveals a high diversification of Y-DNA paternal lineages from haplogroup E: Identifying human male movements within the Mediterranean space. Annals of Human Biology, 2010, 37, 86-107.	0.4	11
26	Revisiting the demographic history of Central African populations from a genetic perspective. , 0, , 1-29.		1