

# Mãrio Vicente

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6760722/publications.pdf>

Version: 2024-02-01

17  
papers

1,376  
citations

758635

12  
h-index

887659

17  
g-index

17  
all docs

17  
docs citations

17  
times ranked

2404  
citing authors

#	ARTICLE	IF	CITATIONS
1	Human adaptation to arsenic in Bolivians living in the Andes. <i>Chemosphere</i> , 2022, 301, 134764.	4.2	7
2	Multiple migrations to the Philippines during the last 50,000 years. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	3.3	50
3	Integrating multi-taxon palaeogenomes and sedimentary ancient DNA to study past ecosystem dynamics. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2021, 288, 20211252.	1.2	14
4	Philippine Ayta possess the highest level of Denisovan ancestry in the world. <i>Current Biology</i> , 2021, 31, 4219-4230.e10.	1.8	37
5	The performance of common SNP arrays in assigning African mitochondrial haplogroups. <i>BMC Genomic Data</i> , 2021, 22, 43.	0.7	2
6	Male-biased migration from East Africa introduced pastoralism into southern Africa. <i>BMC Biology</i> , 2021, 19, 259.	1.7	4
7	Y-Chromosome Variation in Southern African Khoe-San Populations Based on Whole-Genome Sequences. <i>Genome Biology and Evolution</i> , 2020, 12, 1031-1039.	1.1	6
8	Khoe-San Genomes Reveal Unique Variation and Confirm the Deepest Population Divergence in Homo sapiens. <i>Molecular Biology and Evolution</i> , 2020, 37, 2944-2954.	3.5	60
9	African population history: an ancient DNA perspective. <i>Current Opinion in Genetics and Development</i> , 2020, 62, 8-15.	1.5	29
10	Genetic Affinities among Southern Africa Hunter-Gatherers and the Impact of Admixing Farmer and Herder Populations. <i>Molecular Biology and Evolution</i> , 2019, 36, 1849-1861.	3.5	21
11	Population history and genetic adaptation of the Fulani nomads: inferences from genome-wide data and the lactase persistence trait. <i>BMC Genomics</i> , 2019, 20, 915.	1.2	36
12	Population genomics of Mesolithic Scandinavia: Investigating early postglacial migration routes and high-latitude adaptation. <i>PLoS Biology</i> , 2018, 16, e2003703.	2.6	174
13	Southern African ancient genomes estimate modern human divergence to 350,000 to 260,000 years ago. <i>Science</i> , 2017, 358, 652-655.	6.0	351
14	A recent bottleneck of Y chromosome diversity coincides with a global change in culture. <i>Genome Research</i> , 2015, 25, 459-466.	2.4	348
15	A Selective Sweep on a Deleterious Mutation in CPT1A in Arctic Populations. <i>American Journal of Human Genetics</i> , 2014, 95, 584-589.	2.6	119
16	Migration and Interaction in a Contact Zone: mtDNA Variation among Bantu-Speakers in Southern Africa. <i>PLoS ONE</i> , 2014, 9, e99117.	1.1	43
17	Ancient Substructure in Early mtDNA Lineages of Southern Africa. <i>American Journal of Human Genetics</i> , 2013, 92, 285-292.	2.6	75