

Rui Martiniano

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7284392/publications.pdf>

Version: 2024-02-01

24
papers

2,717
citations

471371

17
h-index

580701

25
g-index

33
all docs

33
docs citations

33
times ranked

3633
citing authors

#	ARTICLE	IF	CITATIONS
1	Early farmers from across Europe directly descended from Neolithic Aegeans. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 6886-6891.	3.3	376
2	Upper Palaeolithic genomes reveal deep roots of modern Eurasians. Nature Communications, 2015, 6, 8912.	5.8	334
3	The first horse herders and the impact of early Bronze Age steppe expansions into Asia. Science, 2018, 360, .	6.0	262
4	The population history of northeastern Siberia since the Pleistocene. Nature, 2019, 570, 182-188.	13.7	259
5	Early Neolithic genomes from the eastern Fertile Crescent. Science, 2016, 353, 499-503.	6.0	230
6	Neolithic and Bronze Age migration to Ireland and establishment of the insular Atlantic genome. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 368-373.	3.3	199
7	Ancient genomes indicate population replacement in Early Neolithic Britain. Nature Ecology and Evolution, 2019, 3, 765-771.	3.4	156
8	Dose-dependent expression of claudin-5 is a modifying factor in schizophrenia. Molecular Psychiatry, 2018, 23, 2156-2166.	4.1	148
9	Population genomics of the Viking world. Nature, 2020, 585, 390-396.	13.7	143
10	The population genomics of archaeological transition in west Iberia: Investigation of ancient substructure using imputation and haplotype-based methods. PLoS Genetics, 2017, 13, e1006852.	1.5	122
11	Continuity and Admixture in the Last Five Millennia of Levantine History from Ancient Canaanite and Present-Day Lebanese Genome Sequences. American Journal of Human Genetics, 2017, 101, 274-282.	2.6	102
12	Genomic signals of migration and continuity in Britain before the Anglo-Saxons. Nature Communications, 2016, 7, 10326.	5.8	100
13	Detection of novel germline mutations for breast cancer in non- <i>BRCA1/2</i> families. FEBS Journal, 2015, 282, 3424-3437.	2.2	50
14	Removing reference bias and improving indel calling in ancient DNA data analysis by mapping to a sequence variation graph. Genome Biology, 2020, 21, 250.	3.8	44
15	A Transient Pulse of Genetic Admixture from the Crusaders in the Near East Identified from Ancient Genome Sequences. American Journal of Human Genetics, 2019, 104, 977-984.	2.6	35
16	Insular Celtic population structure and genomic footprints of migration. PLoS Genetics, 2018, 14, e1007152.	1.5	30
17	Genetic Evidence of African Slavery at the Beginning of the Trans-Atlantic Slave Trade. Scientific Reports, 2014, 4, 5994.	1.6	24
18	Placing Ancient DNA Sequences into Reference Phylogenies. Molecular Biology and Evolution, 2022, 39, .	3.5	23

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19	The Eastern side of the Westernmost Europeans: Insights from subclades within Yâ€chromosome haplogroup Jâ€M304. <i>American Journal of Human Biology</i> , 2018, 30, e23082.	0.8	17
20	Ancient DNA at the edge of the world: Continental immigration and the persistence of Neolithic male lineages in Bronze Age Orkney. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, .	3.3	12
21	Biomolecular insights into North African-related ancestry, mobility and diet in eleventh-century Al-Andalus. <i>Scientific Reports</i> , 2021, 11, 18121.	1.6	8
22	Overview of the Americasâ€™ First Peopling from a Patrilineal Perspective: New Evidence from the Southern Continent. <i>Genes</i> , 2022, 13, 220.	1.0	5
23	Y -chromosome diversity in central Portugal reveals signatures of ancient maritime expansions. <i>Anthropologischer Anzeiger</i> , 2013, 70, 355-367.	0.2	3
24	Response to Giam. <i>American Journal of Human Genetics</i> , 2018, 102, 331.	2.6	1