

# 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

471509

17  
h-index

580821

25  
g-index

33  
all docs

33  
docs citations

33  
times ranked

3633  
citing authors

#	ARTICLE	IF	CITATIONS
1	Placing Ancient DNA Sequences into Reference Phylogenies. <i>Molecular Biology and Evolution</i> , 2022, 39, .	8.9	23
2	Overview of the Americasâ€™ First Peopling from a Patrilineal Perspective: New Evidence from the Southern Continent. <i>Genes</i> , 2022, 13, 220.	2.4	5
3	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, .	7.1	12
4	Biomolecular insights into North African-related ancestry, mobility and diet in eleventh-century Al-Andalus. <i>Scientific Reports</i> , 2021, 11, 18121.	3.3	8
5	Removing reference bias and improving indel calling in ancient DNA data analysis by mapping to a sequence variation graph. <i>Genome Biology</i> , 2020, 21, 250.	8.8	44
6	Population genomics of the Viking world. <i>Nature</i> , 2020, 585, 390-396.	27.8	143
7	The population history of northeastern Siberia since the Pleistocene. <i>Nature</i> , 2019, 570, 182-188.	27.8	259
8	A Transient Pulse of Genetic Admixture from the Crusaders in the Near East Identified from Ancient Genome Sequences. <i>American Journal of Human Genetics</i> , 2019, 104, 977-984.	6.2	35
9	Ancient genomes indicate population replacement in Early Neolithic Britain. <i>Nature Ecology and Evolution</i> , 2019, 3, 765-771.	7.8	156
10	Response to G1em. <i>American Journal of Human Genetics</i> , 2018, 102, 331.	6.2	1
11	Dose-dependent expression of claudin-5 is a modifying factor in schizophrenia. <i>Molecular Psychiatry</i> , 2018, 23, 2156-2166.	7.9	148
12	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.	1.6	17
13	The first horse herders and the impact of early Bronze Age steppe expansions into Asia. <i>Science</i> , 2018, 360, .	12.6	262
14	Insular Celtic population structure and genomic footprints of migration. <i>PLoS Genetics</i> , 2018, 14, e1007152.	3.5	30
15	Continuity and Admixture in the Last Five Millennia of Levantine History from Ancient Canaanite and Present-Day Lebanese Genome Sequences. <i>American Journal of Human Genetics</i> , 2017, 101, 274-282.	6.2	102
16	The population genomics of archaeological transition in west Iberia: Investigation of ancient substructure using imputation and haplotype-based methods. <i>PLoS Genetics</i> , 2017, 13, e1006852.	3.5	122
17	Early Neolithic genomes from the eastern Fertile Crescent. <i>Science</i> , 2016, 353, 499-503.	12.6	230
18	Early farmers from across Europe directly descended from Neolithic Aegeans. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 6886-6891.	7.1	376

#	ARTICLE	IF	CITATIONS
19	Genomic signals of migration and continuity in Britain before the Anglo-Saxons. Nature Communications, 2016, 7, 10326.	12.8	100
20	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.	7.1	199
21	Detection of novel germline mutations for breast cancer in non-BRCA1/2 families. FEBS Journal, 2015, 282, 3424-3437.	4.7	50
22	Upper Palaeolithic genomes reveal deep roots of modern Eurasians. Nature Communications, 2015, 6, 8912.	12.8	334
23	Genetic Evidence of African Slavery at the Beginning of the Trans-Atlantic Slave Trade. Scientific Reports, 2014, 4, 5994.	3.3	24
24	Y-chromosome diversity in central Portugal reveals signatures of ancient maritime expansions. Anthropologischer Anzeiger, 2013, 70, 355-367.	0.4	3