

JazmÃ-n Ramos-Madrigal

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3190913/publications.pdf>

Version: 2024-02-01

24
papers

1,492
citations

394421

19
h-index

610901

24
g-index

25
all docs

25
docs citations

25
times ranked

2713
citing authors

#	ARTICLE	IF	CITATIONS
1	Multiproxy evidence highlights a complex evolutionary legacy of maize in South America. <i>Science</i> , 2018, 362, 1309-1313.	12.6	172
2	Next-generation biology: Sequencing and data analysis approaches for non-model organisms. <i>Marine Genomics</i> , 2016, 30, 3-13.	1.1	164
3	Early Pleistocene enamel proteome from Dmanisi resolves <i>Stephanorhinus</i> phylogeny. <i>Nature</i> , 2019, 574, 103-107.	27.8	135
4	Genome Sequence of a 5,310-Year-Old Maize Cob Provides Insights into the Early Stages of Maize Domestication. <i>Current Biology</i> , 2016, 26, 3195-3201.	3.9	130
5	Interspecific Gene Flow Shaped the Evolution of the Genus <i>Canis</i> . <i>Current Biology</i> , 2018, 28, 3441-3449.e5.	3.9	110
6	The dental proteome of <i>Homo antecessor</i> . <i>Nature</i> , 2020, 580, 235-238.	27.8	100
7	Palaeogenomic insights into the origins of French grapevine diversity. <i>Nature Plants</i> , 2019, 5, 595-603.	9.3	85
8	Enamel proteome shows that <i>Gigantopithecus</i> was an early diverging pongine. <i>Nature</i> , 2019, 576, 262-265.	27.8	82
9	Arctic-adapted dogs emerged at the Pleistocene–Holocene transition. <i>Science</i> , 2020, 368, 1495-1499.	12.6	60
10	Near-Random Distribution of Chromosome-Derived Circular DNA in the Condensed Genome of Pigeons and the Larger, More Repeat-Rich Human Genome. <i>Genome Biology and Evolution</i> , 2020, 12, 3762-3777.	2.5	52
11	Grey wolf genomic history reveals a dual ancestry of dogs. <i>Nature</i> , 2022, 607, 313-320.	27.8	48
12	Genomic insights into the early peopling of the Caribbean. <i>Science</i> , 2020, 369, 456-460.	12.6	44
13	New insights on single-stranded versus double-stranded DNA library preparation for ancient DNA. <i>BioTechniques</i> , 2015, 59, 368-371.	1.8	43
14	The limits and potential of paleogenomic techniques for reconstructing grapevine domestication. <i>Journal of Archaeological Science</i> , 2016, 72, 57-70.	2.4	43
15	Genomic Adaptations and Evolutionary History of the Extinct Scimitar-Toothed Cat, <i>Homotherium latidens</i> . <i>Current Biology</i> , 2020, 30, 5018-5025.e5.	3.9	34
16	Tracking the history of grapevine cultivation in Georgia by combining geometric morphometrics and ancient DNA. <i>Vegetation History and Archaeobotany</i> , 2021, 30, 63-76.	2.1	29
17	The genomic landscape of Mexican Indigenous populations brings insights into the peopling of the Americas. <i>Nature Communications</i> , 2021, 12, 5942.	12.8	28
18	An expanded mammal mitogenome dataset from Southeast Asia. <i>GigaScience</i> , 2017, 6, 1-8.	6.4	27

#	ARTICLE	IF	CITATIONS
19	Genomes of Pleistocene Siberian Wolves Uncover Multiple Extinct Wolf Lineages. <i>Current Biology</i> , 2021, 31, 198-206.e8.	3.9	26
20	The efficacy of whole human genome capture on ancient dental calculus and dentin. <i>American Journal of Physical Anthropology</i> , 2019, 168, 496-509.	2.1	24
21	Modern Siberian dog ancestry was shaped by several thousand years of Eurasian-wide trade and human dispersal. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	19
22	Ancient DNA analysis of a nineteenth century tobacco pipe from a Maryland slave quarter. <i>Journal of Archaeological Science</i> , 2019, 105, 11-18.	2.4	15
23	Extended survival of Pleistocene Siberian wolves into the early 20th century on the island of Honshū. <i>IScience</i> , 2021, 24, 101904.	4.1	9
24	Kouprey (<i>Bos sauveli</i>) genomes unveil polytomic origin of wild Asian Bos. <i>IScience</i> , 2021, 24, 103226.	4.1	8