

Cristina E Valdiosera

List of Publications by Year in descending order

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Version: 2024-02-01

28
papers

4,523
citations

279487

23
h-index

476904

29
g-index

33
all docs

33
docs citations

33
times ranked

6650
citing authors

#	ARTICLE	IF	CITATIONS
1	Bioarchaeological evidence of one of the earliest Islamic burials in the Levant. <i>Communications Biology</i> , 2022, 5, .	2.0	3
2	Ancient pigs reveal a near-complete genomic turnover following their introduction to Europe. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 17231-17238.	3.3	101
3	Unraveling the genetic history of the European wild goats. <i>Quaternary Science Reviews</i> , 2018, 185, 189-198.	1.4	21
4	Four millennia of Iberian biomolecular prehistory illustrate the impact of prehistoric migrations at the far end of Eurasia. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 3428-3433.	3.3	96
5	Population genomics of Mesolithic Scandinavia: Investigating early postglacial migration routes and high-latitude adaptation. <i>PLoS Biology</i> , 2018, 16, e2003703.	2.6	174
6	An unusual Pre-bell beaker copper age cave burial context from El Portal ³ⁿ de Cueva Mayor site (Sierra de Atapuerca, Burgos). <i>Quaternary International</i> , 2017, 433, 142-155.	0.7	19
7	Comparing ancient DNA survival and proteome content in 69 archaeological cattle tooth and bone samples from multiple European sites. <i>Journal of Proteomics</i> , 2017, 158, 1-8.	1.2	54
8	High-resolution palaeodietary reconstruction: Amino acid $\delta^{13}C$ analysis of keratin from single hairs of mummified human individuals. <i>Quaternary International</i> , 2017, 436, 96-113.	0.7	22
9	Ancient individuals from the North American Northwest Coast reveal 10,000 years of regional genetic continuity. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 4093-4098.	3.3	100
10	Genomic Analyses of Pre-European Conquest Human Remains from the Canary Islands Reveal Close Affinity to Modern North Africans. <i>Current Biology</i> , 2017, 27, 3396-3402.e5.	1.8	62
11	Ancient <i>DNA</i> reveals past existence of <i>Eurasian lynx</i> in <i>Spain</i> . <i>Journal of Zoology</i> , 2016, 298, 94-102.	0.8	11
12	Ancient mitochondrial DNA provides high-resolution time scale of the peopling of the Americas. <i>Science Advances</i> , 2016, 2, e1501385.	4.7	306
13	Genomic Evidence Establishes Anatolia as the Source of the European Neolithic Gene Pool. <i>Current Biology</i> , 2016, 26, 270-275.	1.8	111
14	Ancient DNA evidence of Iberian lynx palaeoendemism. <i>Quaternary Science Reviews</i> , 2015, 112, 172-180.	1.4	25
15	Genomic evidence for the Pleistocene and recent population history of Native Americans. <i>Science</i> , 2015, 349, aab3884.	6.0	449
16	The ancestry and affiliations of Kennewick Man. <i>Nature</i> , 2015, 523, 455-458.	13.7	241
17	Ancient genomes link early farmers from Atapuerca in Spain to modern-day Basques. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 11917-11922.	3.3	255
18	<i>bammds</i> : a tool for assessing the ancestry of low-depth whole-genome data using multidimensional scaling (MDS). <i>Bioinformatics</i> , 2014, 30, 2962-2964.	1.8	40

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19	Genomic Diversity and Admixture Differs for Stone-Age Scandinavian Foragers and Farmers. <i>Science</i> , 2014, 344, 747-750.	6.0	315
20	Pulling out the 1%: Whole-Genome Capture for the Targeted Enrichment of Ancient DNA Sequencing Libraries. <i>American Journal of Human Genetics</i> , 2013, 93, 852-864.	2.6	284
21	Complete mitochondrial genome sequence of a Middle Pleistocene cave bear reconstructed from ultrashort DNA fragments. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 15758-15763.	3.3	1,097
22	Ancient mtDNA identifies postglacial recolonisation, not recent bottlenecks, as the primary driver of contemporary mtDNA phylogeography and diversity in Scandinavian brown bears. <i>Diversity and Distributions</i> , 2013, 19, 245-256.	1.9	59
23	50,000 years of genetic uniformity in the critically endangered Iberian lynx. <i>Molecular Ecology</i> , 2011, 20, 3785-3795.	2.0	30
24	Ancient Hybridization and an Irish Origin for the Modern Polar Bear Matriline. <i>Current Biology</i> , 2011, 21, 1251-1258.	1.8	257
25	Surprising migration and population size dynamics in ancient Iberian brown bears (<i>Ursus arctos</i>). <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 5123-5128.	3.3	86
26	Ancient DNA reveals lack of postglacial habitat tracking in the arctic fox. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 6726-6729.	3.3	137
27	Staying out in the cold: glacial refugia and mitochondrial DNA phylogeography in ancient European brown bears. <i>Molecular Ecology</i> , 2007, 16, 5140-5148.	2.0	130
28	Typing single polymorphic nucleotides in mitochondrial DNA as a way to access Middle Pleistocene DNA. <i>Biology Letters</i> , 2006, 2, 601-603.	1.0	28