## Paula F Campos

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7796307/publications.pdf

Version: 2024-02-01

58 papers 9,642 citations

33 h-index 60 g-index

64 all docs

64
docs citations

64 times ranked 13236 citing authors

#	Article	IF	CITATIONS
1	The era of reference genomes in conservation genomics. Trends in Ecology and Evolution, 2022, 37, 197-202.	4.2	138
2	Genetic diversity of the endangered Mongolian saiga antelope <i>Saiga tatarica mongolic</i> a (Artiodactyla: Bovidae) provides insights into conservation. Biological Journal of the Linnean Society, 2022, 137, 100-111.	0.7	4
3	Mitogenomics of the endangered Mediterranean monk seal ( <i>Monachus monachus</i> ) reveals dramatic loss of diversity and supports historical gene-flow between Atlantic and eastern Mediterranean populations. Zoological Journal of the Linnean Society, 2021, 191, 1147-1159.	1.0	8
4	Ancient DNA reveals the lost domestication history of South American camelids in Northern Chile and across the Andes. ELife, $2021,10,10$	2.8	31
5	Exploring the phylogeography and population dynamics of the giant deer ( <i>Megaloceros) Tj ETQq1 1 0.784314 Sciences, 2021, 288, 20201864.</i>	4 rgBT /Ov 1.2	verlock 10 Tf ! 6
6	Comparative genomics provides insights into the aquatic adaptations of mammals. Proceedings of the National Academy of Sciences of the United States of America, $2021,118,.$	3.3	43
7	Response of an Afro-Palearctic bird migrant to glaciation cycles. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	3.3	25
8	Declining genetic diversity of European honeybees along the twentieth century. Scientific Reports, 2020, 10, 10520.	1.6	41
9	Complete Inactivation of Sebum-Producing Genes Parallels the Loss of Sebaceous Glands in Cetacea. Molecular Biology and Evolution, 2019, 36, 1270-1280.	3.5	30
10	DNA Extraction from Keratin and Chitin. Methods in Molecular Biology, 2019, 1963, 57-63.	0.4	7
11	Parallel adaptation of rabbit populations to myxoma virus. Science, 2019, 363, 1319-1326.	6.0	124
12	Genomic insights into the origin and diversification of late maritime hunter-gatherers from the Chilean Patagonia. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E4006-E4012.	3.3	50
13	The genetic history of whaling in the Cantabrian Sea during the 13th–18th centuries: Were North Atlantic right whales (Eubalaena glacialis) the main target species?. Journal of Archaeological Science: Reports, 2018, 18, 393-398.	0.2	2
14	Phylogenetic position of the extinct blue antelope, Hippotragus leucophaeus (Pallas, 1766) (Bovidae:) Tj ETQq0 C 2018, 182, 225-235.	0 0 rgBT /C 1.0	Overlock 10 Tf 9
15	"Out of the Can― A Draft Genome Assembly, Liver Transcriptome, and Nutrigenomics of the European Sardine, Sardina pilchardus. Genes, 2018, 9, 485.	1.0	30
16	Phylogenomics and Morphology of Extinct Paleognaths Reveal the Origin and Evolution of the Ratites. Current Biology, 2017, 27, 68-77.	1.8	123
17	Mitochondrial DNA of preâ€ast glacial maximum red deer from NW Spain suggests a more complex phylogeographical history for the species. Ecology and Evolution, 2017, 7, 10690-10700.	0.8	13
18	Next-generation biology: Sequencing and data analysis approaches for non-model organisms. Marine Genomics, 2016, 30, 3-13.	0.4	164

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19	A genomic history of Aboriginal Australia. Nature, 2016, 538, 207-214.	13.7	439
20	Metagenomic Analysis from the Interior of a Speleothem in Tjuv-Ante's Cave, Northern Sweden. PLoS ONE, 2016, 11, e0151577.	1.1	29
21	Interordinal gene capture, the phylogenetic position of Steller's sea cow based on molecular and morphological data, and the macroevolutionary history of Sirenia. Molecular Phylogenetics and Evolution, 2015, 91, 178-193.	1.2	75
22	Ancient genomics. Philosophical Transactions of the Royal Society B: Biological Sciences, 2015, 370, 20130387.	1.8	142
23	Genomic evidence for the Pleistocene and recent population history of Native Americans. Science, 2015, 349, aab3884.	6.0	449
24	Complete mitochondrial DNA sequence of the endangered giant sable antelope (Hippotragus niger) Tj ETQq0 0 242-249.	0 rgBT /Ov 1.2	verlock 10 Tf
25	Ancient and modern environmental DNA. Philosophical Transactions of the Royal Society B: Biological Sciences, 2015, 370, 20130383.	1.8	292
26	Special report: Anatomical pathology A glimpse into the early origins of medieval anatomy through the oldest conserved human dissection (Western Europe, 13th c. A.D.). Archives of Medical Science, 2014, 2, 366-373.	0.4	13
27	Whole-genome analyses resolve early branches in the tree of life of modern birds. Science, 2014, 346, 1320-1331.	6.0	1,583
28	Comparative genomics reveals insights into avian genome evolution and adaptation. Science, 2014, 346, 1311-1320.	6.0	895
29	Upper Palaeolithic Siberian genome reveals dual ancestry of Native Americans. Nature, 2014, 505, 87-91.	13.7	821
30	Two ancient human genomes reveal Polynesian ancestry among the indigenous Botocudos of Brazil. Current Biology, 2014, 24, R1035-R1037.	1.8	73
31	Musk ox (Ovibos moschatus) of the mammoth steppe: tracing palaeodietary and palaeoenvironmental changes over the last 50,000 years using carbon and nitrogen isotopic analysis. Quaternary Science Reviews, 2014, 102, 192-201.	1.4	27
32	Colonization history of Mallorca Island by the European rabbit, <i>Oryctolagus cuniculus </i> , and the Iberian hare, <i>Lepus granatensis </i> (Lagomorpha: Leporidae). Biological Journal of the Linnean Society, 2014, 111, 748-760.	0.7	7
33	Identification of Polynesian mtDNA haplogroups in remains of Botocudo Amerindians from Brazil. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 6465-6469.	3.3	42
34	Mitochondrial genome diversity and population structure of the giant squid <i>Architeuthis</i> genetics sheds new light on one of the most enigmatic marine species. Proceedings of the Royal Society B: Biological Sciences, 2013, 280, 20130273.	1.2	57
35	Independent confirmation of a diagnostic sheep/goat peptide sequence through DNA analysis and further exploration of its taxonomic utility within the Bovidae. Journal of Archaeological Science, 2013, 40, 1421-1424.	1.2	11
36	Reconstructing genome evolution in historic samples of the Irish potato famine pathogen. Nature Communications, 2013, 4, 2172.	5.8	103

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37	Clovis Age Western Stemmed Projectile Points and Human Coprolites at the Paisley Caves. Science, 2012, 337, 223-228.	6.0	211
38	DNA Extraction from Formalin-Fixed Material. Methods in Molecular Biology, 2012, 840, 81-85.	0.4	80
39	The half-life of DNA in bone: measuring decay kinetics in 158 dated fossils. Proceedings of the Royal Society B: Biological Sciences, 2012, 279, 4724-4733.	1.2	478
40	DNA Extraction from Keratin and Chitin. Methods in Molecular Biology, 2012, 840, 43-49.	0.4	20
41	DNA in ancient bone – Where is it located and how should we extract it?. Annals of Anatomy, 2012, 194, 7-16.	1.0	132
42	Species-specific responses of Late Quaternary megafauna to climate and humans. Nature, 2011, 479, 359-364.	13.7	586
43	Genetic evidence for patrilocal mating behavior among Neandertal groups. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 250-253.	3.3	165
44	A multidisciplinary study of archaeological grape seeds. Die Naturwissenschaften, 2010, 97, 205-217.	0.6	82
45	Molecular identification of the extinct mountain goat, <i>Oreamnos harringtoni</i> (Bovidae). Boreas, 2010, 39, 18-23.	1.2	5
46	Ancient DNA sequences point to a large loss of mitochondrial genetic diversity in the saiga antelope ( <i>Saiga tatarica</i> ) since the Pleistocene. Molecular Ecology, 2010, 19, 4863-4875.	2.0	59
47	Ancient human genome sequence of an extinct Palaeo-Eskimo. Nature, 2010, 463, 757-762.	13.7	750
48	Ancient DNA analyses exclude humans as the driving force behind late Pleistocene musk ox ( <i>Ovibos) Tj ETQq0 States of America, 2010, 107, 5675-5680.</i>	0 0 0 rgBT 3.3	Overlock 10 208
49	Multidisciplinary medical identification of a French king's head (Henri IV). BMJ, The, 2010, 341, c6805-c6805.	3.0	24
50	Ancient DNA, a Neolithic legging from the Swiss Alps and the early history of goat. Journal of Archaeological Science, 2010, 37, 1247-1251.	1.2	27
51	Clarification of the taxonomic relationship of the extant and extinct ovibovids, Ovibos, Praeovibos, Euceratherium and Bootherium. Quaternary Science Reviews, 2010, 29, 2123-2130.	1.4	17
52	Analysis of complete mitochondrial genomes from extinct and extant rhinoceroses reveals lack of phylogenetic resolution. BMC Evolutionary Biology, 2009, 9, 95.	3.2	92
53	Paleo-Eskimo mtDNA Genome Reveals Matrilineal Discontinuity in Greenland. Science, 2008, 320, 1787-1789.	6.0	184
54	Intraspecific phylogenetic analysis of Siberian woolly mammoths using complete mitochondrial genomes. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 8327-8332.	3.3	149

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55	Historical Mammal Extinction on Christmas Island (Indian Ocean) Correlates with Introduced Infectious Disease. PLoS ONE, 2008, 3, e3602.	1.1	198
56	Whole-Genome Shotgun Sequencing of Mitochondria from Ancient Hair Shafts. Science, 2007, 317, 1927-1930.	6.0	220
57	5′-Tailed sequencing primers improve sequencing quality of PCR products. BioTechniques, 2007, 42, 174-176.	0.8	33
58	Comparative Analyses of 35 Marine Mammal Genomes Provide Insights into the Evolution of Aquatic Life. SSRN Electronic Journal, 0, , .	0.4	0