

# Paula F Campos

## List of Publications by Year in descending order

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Version: 2024-02-01

58  
papers

9,642  
citations

126708

33  
h-index

128067

60  
g-index

64  
all docs

64  
docs citations

64  
times ranked

13236  
citing authors



#	ARTICLE	IF	CITATIONS
19	A genomic history of Aboriginal Australia. <i>Nature</i> , 2016, 538, 207-214.	13.7	439
20	Metagenomic Analysis from the Interior of a Speleothem in Tjuv-Ante's Cave, Northern Sweden. <i>PLoS ONE</i> , 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. <i>Molecular Phylogenetics and Evolution</i> , 2015, 91, 178-193.	1.2	75
22	Ancient genomics. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2015, 370, 20130387.	1.8	142
23	Genomic evidence for the Pleistocene and recent population history of Native Americans. <i>Science</i> , 2015, 349, aab3884.	6.0	449
24	Complete mitochondrial DNA sequence of the endangered giant sable antelope ( <i>Hippotragus niger</i> ) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5 242-249.	1.2	9
25	Ancient and modern environmental DNA. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 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.). <i>Archives of Medical Science</i> , 2014, 2, 366-373.	0.4	13
27	Whole-genome analyses resolve early branches in the tree of life of modern birds. <i>Science</i> , 2014, 346, 1320-1331.	6.0	1,583
28	Comparative genomics reveals insights into avian genome evolution and adaptation. <i>Science</i> , 2014, 346, 1311-1320.	6.0	895
29	Upper Palaeolithic Siberian genome reveals dual ancestry of Native Americans. <i>Nature</i> , 2014, 505, 87-91.	13.7	821
30	Two ancient human genomes reveal Polynesian ancestry among the indigenous Botocudos of Brazil. <i>Current Biology</i> , 2014, 24, R1035-R1037.	1.8	73
31	Musk ox ( <i>Ovibos moschatus</i> ) of the mammoth steppe: tracing palaeodietary and palaeoenvironmental changes over the last 50,000 years using carbon and nitrogen isotopic analysis. <i>Quaternary Science Reviews</i> , 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). <i>Biological Journal of the Linnean Society</i> , 2014, 111, 748-760.	0.7	7
33	Identification of Polynesian mtDNA haplogroups in remains of Botocudo Amerindians from Brazil. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 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. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 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. <i>Journal of Archaeological Science</i> , 2013, 40, 1421-1424.	1.2	11
36	Reconstructing genome evolution in historic samples of the Irish potato famine pathogen. <i>Nature Communications</i> , 2013, 4, 2172.	5.8	103

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37	Clovis Age Western Stemmed Projectile Points and Human Coprolites at the Paisley Caves. <i>Science</i> , 2012, 337, 223-228.	6.0	211
38	DNA Extraction from Formalin-Fixed Material. <i>Methods in Molecular Biology</i> , 2012, 840, 81-85.	0.4	80
39	The half-life of DNA in bone: measuring decay kinetics in 158 dated fossils. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2012, 279, 4724-4733.	1.2	478
40	DNA Extraction from Keratin and Chitin. <i>Methods in Molecular Biology</i> , 2012, 840, 43-49.	0.4	20
41	DNA in ancient bone – Where is it located and how should we extract it?. <i>Annals of Anatomy</i> , 2012, 194, 7-16.	1.0	132
42	Species-specific responses of Late Quaternary megafauna to climate and humans. <i>Nature</i> , 2011, 479, 359-364.	13.7	586
43	Genetic evidence for patrilocal mating behavior among Neandertal groups. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 250-253.	3.3	165
44	A multidisciplinary study of archaeological grape seeds. <i>Die Naturwissenschaften</i> , 2010, 97, 205-217.	0.6	82
45	Molecular identification of the extinct mountain goat, <i>Oreamnos harringtoni</i> (Bovidae). <i>Boreas</i> , 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. <i>Molecular Ecology</i> , 2010, 19, 4863-4875.	2.0	59
47	Ancient human genome sequence of an extinct Palaeo-Eskimo. <i>Nature</i> , 2010, 463, 757-762.	13.7	750
48	Ancient DNA analyses exclude humans as the driving force behind late Pleistocene musk ox ( <i>Ovibos</i> ) in the States of America, 2010, 107, 5675-5680.	3.3	208
49	Multidisciplinary medical identification of a French king's head (Henri IV). <i>BMJ</i> , 2010, 341, c6805-c6805.	3.0	24
50	Ancient DNA, a Neolithic legging from the Swiss Alps and the early history of goat. <i>Journal of Archaeological Science</i> , 2010, 37, 1247-1251.	1.2	27
51	Clarification of the taxonomic relationship of the extant and extinct ovibovids, <i>Ovibos</i> , <i>Praeovibos</i> , <i>Euceratherium</i> and <i>Bootherium</i> . <i>Quaternary Science Reviews</i> , 2010, 29, 2123-2130.	1.4	17
52	Analysis of complete mitochondrial genomes from extinct and extant rhinoceroses reveals lack of phylogenetic resolution. <i>BMC Evolutionary Biology</i> , 2009, 9, 95.	3.2	92
53	Paleo-Eskimo mtDNA Genome Reveals Matrilineal Discontinuity in Greenland. <i>Science</i> , 2008, 320, 1787-1789.	6.0	184
54	Intraspecific phylogenetic analysis of Siberian woolly mammoths using complete mitochondrial genomes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 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'â€²-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