

Charlotte Lindqvist

List of Publications by Year in descending order

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Version: 2024-02-01

29
papers

1,441
citations

430442

18
h-index

500791

28
g-index

31
all docs

31
docs citations

31
times ranked

2163
citing authors

#	ARTICLE	IF	CITATIONS
1	Polar and brown bear genomes reveal ancient admixture and demographic footprints of past climate change. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, E2382-90.	3.3	310
2	Complete mitochondrial genome of a Pleistocene jawbone unveils the origin of polar bear. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 5053-5057.	3.3	132
3	Origin of the Hawaiian endemic mints within North American <i>Stachys</i> (Lamiaceae). <i>American Journal of Botany</i> , 2002, 89, 1709-1724.	0.8	125
4	An updated phylogeny and classification of Lamiaceae subfamily Lamioideae. <i>Taxon</i> , 2011, 60, 471-484.	0.4	122
5	Deglaciation of the Pacific coastal corridor directly preceded the human colonization of the Americas. <i>Science Advances</i> , 2018, 4, eaar5040.	4.7	107
6	Molecular Phylogenetics, Character Evolution, and Suprageneric Classification of Lamioideae (Lamiaceae). <i>Annals of the Missouri Botanical Garden</i> , 2010, 97, 191-217.	1.3	81
7	Evolutionary history of enigmatic bears in the Tibetan Plateau-Himalaya region and the identity of the yeti. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2017, 284, 20171804.	1.2	62
8	Molecular phylogenetics of an aquatic plant lineage, Potamogetonaceae. <i>Cladistics</i> , 2006, 22, 568-588.	1.5	61
9	Polar Bears Exhibit Genome-Wide Signatures of Bioenergetic Adaptation to Life in the Arctic Environment. <i>Genome Biology and Evolution</i> , 2014, 6, 433-450.	1.1	52
10	Cladogenesis and reticulation in the Hawaiian endemic mints (Lamiaceae). <i>Cladistics</i> , 2003, 19, 480-495.	1.5	50
11	The quest to resolve recent radiations: Plastid phylogenomics of extinct and endangered Hawaiian endemic mints (Lamiaceae). <i>Molecular Phylogenetics and Evolution</i> , 2016, 99, 16-33.	1.2	47
12	Molecular phylogeny of tribe Stachydeae (Lamiaceae subfamily Lamioideae). <i>Molecular Phylogenetics and Evolution</i> , 2013, 69, 535-551.	1.2	40
13	Phylogeny and biogeography of New World Stachydeae (Lamiaceae) with emphasis on the origin and diversification of Hawaiian and South American taxa. <i>Molecular Phylogenetics and Evolution</i> , 2013, 69, 218-238.	1.2	37
14	An expressed sequence tag (EST) library from developing fruits of an Hawaiian endemic mint (<i>Stenogyne rugosa</i> , Lamiaceae): characterization and microsatellite markers. <i>BMC Plant Biology</i> , 2006, 6, 16.	1.6	30
15	Untangling reticulate evolutionary relationships among New World and Hawaiian mints (Stachydeae.) <i>Tj ETQq1 1 0,784314 rgBT /Over</i>	1.2	30
16	Whole-genome analysis of <i>Mustela erminea</i> finds that pulsed hybridization impacts evolution at high latitudes. <i>Communications Biology</i> , 2018, 1, 51.	2.0	24
17	Polyglutamine variation in a flowering time protein correlates with island age in a Hawaiian plant radiation. <i>BMC Evolutionary Biology</i> , 2007, 7, 105.	3.2	22
18	Molecular phylogenetics of tribe Synandreae, a North American lineage of lamioid mints (Lamiaceae). <i>Cladistics</i> , 2008, 24, 299-314.	1.5	18

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19	New insights into evolutionary relationships within the subfamily Lamiioideae (Lamiaceae) based on pentatricopeptide repeat (<i>PPR</i>) nuclear DNA sequences. American Journal of Botany, 2015, 102, 1721-1735.	0.8	18
20	An early dog from southeast Alaska supports a coastal route for the first dog migration into the Americas. Proceedings of the Royal Society B: Biological Sciences, 2021, 288, 20203103.	1.2	17
21	Insights into bear evolution from a Pleistocene polar bear genome. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, .	3.3	11
22	Genetic diversity of historical Atlantic walruses (<i>Odobenus rosmarus rosmarus</i>) from Bj�rn�ya and H�y�ya (Tusen�yane), Svalbard, Norway. BMC Research Notes, 2016, 9, 112.	0.6	10
23	Whole�genome resequencing reveals persistence of forest�associated mammals in Late Pleistocene refugia along North America�s North Pacific Coast. Journal of Biogeography, 2021, 48, 1153-1169.	1.4	7
24	Mitochondrial DNA variation of a natural population of <i>Gyrodactylus thymalli</i> (Monogenea) from the type locality River Hnilec, Slovakia. Parasitology Research, 2007, 101, 1439-1442.	0.6	4
25	Data characterizing the chloroplast genomes of extinct and endangered Hawaiian endemic mints (Lamiaceae) and their close relatives. Data in Brief, 2016, 7, 900-922.	0.5	4
26	Paleogenomics: Genome-Scale Analysis of Ancient DNA and Population and Evolutionary Genomic Inferences. Population Genomics, 2018, , 323-360.	0.2	4
27	Evolutionary relationships within the lamioid tribe Synandreae (Lamiaceae) based on multiple low-copy nuclear loci. PeerJ, 2016, 4, e2220.	0.9	4
28	Palaeoecological and genetic analyses of Late Pleistocene bears in Asiatic Russia. Boreas, 2022, 51, 465-480.	1.2	3
29	Technical Advances and Challenges in Genome-Scale Analysis of Ancient DNA. Population Genomics, 2018, , 3-29.	0.2	2