

Cornelya F C KlÃ¼tsch

List of Publications by Year in descending order

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34
papers

1,173
citations

567281

15
h-index

395702

33
g-index

36
all docs

36
docs citations

36
times ranked

1756
citing authors

#	ARTICLE	IF	CITATIONS
1	mtDNA Data Indicate a Single Origin for Dogs South of Yangtze River, Less Than 16,300 Years Ago, from Numerous Wolves. <i>Molecular Biology and Evolution</i> , 2009, 26, 2849-2864.	8.9	314
2	Mitochondrial DNA data indicate an introduction through Mainland Southeast Asia for Australian dingoes and Polynesian domestic dogs. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2012, 279, 967-974.	2.6	110
3	Mitochondrial and nuclear phylogeny of circum-Mediterranean tree frogs from the <i>Hyla arborea</i> group. <i>Molecular Phylogenetics and Evolution</i> , 2008, 49, 1019-1024.	2.7	93
4	Phylogeography of the Middle Eastern tree frogs (<i>Hyla</i> , Hylidae, Amphibia) as inferred from nuclear and mitochondrial DNA variation, with a description of a new species. <i>Molecular Phylogenetics and Evolution</i> , 2010, 55, 1146-1166.	2.7	92
5	Molecular taxonomy in pholcid spiders (Pholcidae, Araneae): evaluation of species identification methods using CO1 and 16S rRNA. <i>Zoologica Scripta</i> , 2006, 35, 441-457.	1.7	66
6	Comprehensive study of mtDNA among Southwest Asian dogs contradicts independent domestication of wolf, but implies dog-wolf hybridization. <i>Ecology and Evolution</i> , 2011, 1, 373-385.	1.9	59
7	Pre-Columbian origins of Native American dog breeds, with only limited replacement by European dogs, confirmed by mtDNA analysis. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2013, 280, 20131142.	2.6	46
8	Łeghágots'enetę (learning together): the importance of indigenous perspectives in the identification of biological variation. <i>Ecology and Society</i> , 2016, 21, .	2.3	43
9	Phylogeographical Analysis of mtDNA Data Indicates Postglacial Expansion from Multiple Glacial Refugia in Woodland Caribou (<i>Rangifer tarandus caribou</i>). <i>PLoS ONE</i> , 2012, 7, e52661.	2.5	40
10	Conservation genomics in perspective: A holistic approach to understanding <i>Canis</i> evolution in North America. <i>Biological Conservation</i> , 2012, 155, 186-192.	4.1	37
11	Regional occurrence, high frequency but low diversity of mitochondrial DNA haplogroup d1 suggests a recent dog-wolf hybridization in Scandinavia. <i>Animal Genetics</i> , 2011, 42, 100-103.	1.7	32
12	Ancient diversification in glacial refugia leads to intraspecific diversity in a Holarctic mammal. <i>Journal of Biogeography</i> , 2017, 44, 386-396.	3.0	28
13	The eastern migratory caribou: the role of genetic introgression in ecotype evolution. <i>Royal Society Open Science</i> , 2016, 3, 150469.	2.4	27
14	Genetic changes caused by restocking and hydroelectric dams in demographically bottlenecked brown trout in a transnational subarctic riverine system. <i>Ecology and Evolution</i> , 2019, 9, 6068-6081.	1.9	19
15	Population dynamics of caribou shaped by glacial cycles before the last glacial maximum. <i>Molecular Ecology</i> , 2021, 30, 6121-6143.	3.9	19
16	Spatial differences in genetic diversity and northward migration suggest genetic erosion along the boreal caribou southern range limit and continued range retraction. <i>Ecology and Evolution</i> , 2019, 9, 7030-7046.	1.9	16
17	Sea ice reduction drives genetic differentiation among Barents Sea polar bears. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2021, 288, 20211741.	2.6	15
18	Evolutionary reconstruction supports the presence of a Pleistocene Arctic refugium for a large mammal species. <i>Journal of Biogeography</i> , 2017, 44, 2729-2739.	3.0	14

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19	Genetic and morphometric differentiation among island populations of two <i>Norops</i> lizards (Reptilia: Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 622 Td Biogeography, 2007, 34, 1124-1135.	3.0	11
20	Genetic analysis indicates spatial-dependent patterns of sex-biased dispersal in Eurasian lynx in Finland. PLoS ONE, 2021, 16, e0246833.	2.5	11
21	Segregation of point mutation heteroplasmy in the control region of dog mtDNA studied systematically in deep generation pedigrees. International Journal of Legal Medicine, 2011, 125, 527-535.	2.2	10
22	Improved genotyping and sequencing success rates for North American river otter (<i>Lontra</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 622 Td	1.4	9
23	Parallel evolution of site-specific changes in divergent caribou lineages. Ecology and Evolution, 2018, 8, 6053-6064.	1.9	9
24	Combining multiple analytical approaches for the identification of population structure and genetic delineation of two subspecies of the endemic Arabian burnet moth <i>Reissita simonyi</i> (Zygaenidae;) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 622 Td	1.4	9
25	Hybridization of domestic mink with wild American mink (<i>Neovison vison</i>) in eastern Canada. Canadian Journal of Zoology, 2017, 95, 443-451.	1.0	8
26	The IGF1 small dog haplotype is derived from Middle Eastern grey wolves: a closer look at statistics, sampling, and the alleged Middle Eastern origin of small dogs. BMC Biology, 2010, 8, 119.	3.8	6
27	Large-scale genetic admixture suggests high dispersal in an insect pest, the apple fruit moth. PLoS ONE, 2020, 15, e0236509.	2.5	5
28	Distribution of the Yellow-lemon Tree Frog, <i>Hyla savignyi</i> (Audouin, 1827), in southern Arabia: updates and extensions of previous records. Zoology in the Middle East, 2004, 31, 47-52.	0.6	4
29	On tree frog cryptozoology and systematics – response to Y. Werner. Molecular Phylogenetics and Evolution, 2010, 57, 957-958.	2.7	4
30	Does connectivity exist for remnant boreal caribou (<i>Rangifer tarandus caribou</i>) along the Lake Superior Coastal Range? Options for landscape restoration. Rangifer, 2018, 38, 13-26.	0.6	4
31	Characterization of microsatellite loci for <i>Reissita simonyi</i> (Rebel, 1899) (Lepidoptera, Zygaenidae). Molecular Ecology Notes, 2003, 3, 528-531.	1.7	3
32	Studying phenological phenomena in subarctic biomes with international school pupils as citizen scientists. Ecology and Evolution, 2021, 11, 3501-3515.	1.9	3
33	Mitogenomics of the suborder Cottoidei (Teleostei: Perciformes): Improved assemblies, mitogenome features, phylogeny, and ecological implications. Genomics, 2022, 114, 110297.	2.9	3
34	Evaluating evolutionary history and adaptive differentiation to identify conservation units of Canada lynx (<i>Lynx canadensis</i>). Global Ecology and Conservation, 2019, 20, e00708.	2.1	1