

Christian Roos

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7271748/publications.pdf>

Version: 2024-02-01

136
papers

7,575
citations

66343

42
h-index

62596

80
g-index

144
all docs

144
docs citations

144
times ranked

7360
citing authors

#	ARTICLE	IF	CITATIONS
1	A Molecular Phylogeny of Living Primates. PLoS Genetics, 2011, 7, e1001342.	3.5	1,130
2	Impending extinction crisis of the world's primates: Why primates matter. Science Advances, 2017, 3, e1600946.	10.3	912
3	Gibbon genome and the fast karyotype evolution of small apes. Nature, 2014, 513, 195-201.	27.8	320
4	Primate jumping genes elucidate strepsirrhine phylogeny. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 10650-10654.	7.1	238
5	A Mitogenomic Phylogeny of Living Primates. PLoS ONE, 2013, 8, e69504.	2.5	217
6	Sustained virologic control in SIV macaques after antiretroviral and $\hat{\pm}$ antibody therapy. Science, 2016, 354, 197-202.	12.6	194
7	Morphometric, Behavioral, and Genomic Evidence for a New Orangutan Species. Current Biology, 2017, 27, 3487-3498.e10.	3.9	192
8	Whole-genome sequencing of the snub-nosed monkey provides insights into folivory and evolutionary history. Nature Genetics, 2014, 46, 1303-1310.	21.4	174
9	Mitochondrial phylogeography of baboons (Papio spp.) – Indication for introgressive hybridization?. BMC Evolutionary Biology, 2009, 9, 83.	3.2	173
10	The strange blood: Natural hybridization in primates. Evolutionary Anthropology, 2011, 20, 96-103.	3.4	146
11	Nuclear versus mitochondrial DNA: evidence for hybridization in colobine monkeys. BMC Evolutionary Biology, 2011, 11, 77.	3.2	123
12	Primates in peril: the significance of Brazil, Madagascar, Indonesia and the Democratic Republic of the Congo for global primate conservation. PeerJ, 2018, 6, e4869.	2.0	123
13	The comparative genomics and complex population history of <i>Papio</i> baboons. Science Advances, 2019, 5, eaau6947.	10.3	115
14	Mitochondrial evidence for multiple radiations in the evolutionary history of small apes. BMC Evolutionary Biology, 2010, 10, 74.	3.2	111
15	Baboon phylogeny as inferred from complete mitochondrial genomes. American Journal of Physical Anthropology, 2013, 150, 133-140.	2.1	110
16	Molecular Phylogeny of the Major Hylobatid Divisions. Molecular Phylogenetics and Evolution, 2001, 19, 486-494.	2.7	103
17	Phylogenetic position of the langur genera Semnopithecus and Trachypithecus among Asian colobines, and genus affiliations of their species groups. BMC Evolutionary Biology, 2008, 8, 58.	3.2	94
18	Molecular phylogeny and evolutionary history of Southeast Asian macaques forming the <i>M. silenus</i> group. Molecular Phylogenetics and Evolution, 2007, 42, 807-816.	2.7	89

#	ARTICLE	IF	CITATIONS
19	Mitochondrial Diversity and Distribution of African Green Monkeys (<i>Chlorocebus</i> Gray, 1870). <i>American Journal of Primatology</i> , 2013, 75, 350-360.	1.7	87
20	Description of a new species of <i>Hoolock</i> gibbon (Primates: Hylobatidae) based on integrative taxonomy. <i>American Journal of Primatology</i> , 2017, 79, e22631.	1.7	80
21	Concordance between vocal and genetic diversity in crested gibbons. <i>BMC Evolutionary Biology</i> , 2011, 11, 36.	3.2	75
22	Mitochondrial Genome Sequences Effectively Reveal the Phylogeny of <i>Hylobates</i> Gibbons. <i>PLoS ONE</i> , 2010, 5, e14419.	2.5	75
23	Remarkable ancient divergences amongst neglected loriform primates. <i>Zoological Journal of the Linnean Society</i> , 2015, 175, 661-674.	2.3	71
24	Primate phylogeny: molecular evidence from retroposons. <i>Cytogenetic and Genome Research</i> , 2005, 108, 26-37.	1.1	66
25	Is the New Primate Genus <i>Rungwecebus</i> a Baboon?. <i>PLoS ONE</i> , 2009, 4, e4859.	2.5	66
26	Mitochondrial phylogeny of tamarins (<i>Saguinus</i> , Hoffmannsegg 1807) with taxonomic and biogeographic implications for the <i>S. nigricollis</i> species group. <i>American Journal of Physical Anthropology</i> , 2011, 144, 564-574.	2.1	65
27	Retropositional events consolidate the branching order among New World monkey genera. <i>Molecular Phylogenetics and Evolution</i> , 2009, 50, 507-513.	2.7	60
28	Molecular phylogeny and taxonomic revision of the sportive lemurs (<i>Lepilemur</i> , Primates). <i>BMC Evolutionary Biology</i> , 2006, 6, 17.	3.2	59
29	Taxonomic review of the New World tamarins (Primates: Callitrichidae). <i>Zoological Journal of the Linnean Society</i> , 2016, 177, 1003-1028.	2.3	59
30	Species definitions and conservation: a review and case studies from African mammals. <i>Conservation Genetics</i> , 2017, 18, 1247-1256.	1.5	58
31	Hybridization in human evolution: Insights from other organisms. <i>Evolutionary Anthropology</i> , 2019, 28, 189-209.	3.4	57
32	Mitogenomic phylogeny of the common long-tailed macaque (<i>Macaca fascicularis fascicularis</i>). <i>BMC Genomics</i> , 2015, 16, 222.	2.8	55
33	Genome typing of nonhuman primate models: implications for biomedical research. <i>Trends in Genetics</i> , 2014, 30, 482-487.	6.7	54
34	Mitochondrial phylogeny, taxonomy and biogeography of the silvered langur species group (<i>Trachypithecus cristatus</i>). <i>Molecular Phylogenetics and Evolution</i> , 2008, 47, 629-636.	2.7	53
35	Evolutionary History of the Odd-Nosed Monkeys and the Phylogenetic Position of the Newly Described Myanmar Snub-Nosed Monkey <i>Rhinopithecus strykeri</i> . <i>PLoS ONE</i> , 2012, 7, e37418.	2.5	53
36	Introgressive hybridization in southern African baboons shapes patterns of mtDNA variation. <i>American Journal of Physical Anthropology</i> , 2010, 142, 125-136.	2.1	52

#	ARTICLE	IF	CITATIONS
37	A Severe Lack of Evidence Limits Effective Conservation of the World's Primates. <i>BioScience</i> , 2020, 70, 794-803.	4.9	51
38	The rise and fall of a genus: Complete mtDNA genomes shed light on the phylogenetic position of yellow-tailed woolly monkeys, <i>Lagothrix flavicauda</i> , and on the evolutionary history of the family Atelidae (Primates: Platyrrhini). <i>Molecular Phylogenetics and Evolution</i> , 2015, 82, 495-510.	2.7	50
39	Mitochondrial phylogeny of leaf monkeys (genus <i>Presbytis</i> , Eschscholtz, 1821) with implications for taxonomy and conservation. <i>Molecular Phylogenetics and Evolution</i> , 2011, 59, 311-319.	2.7	49
40	Mitogenomics of the Old World monkey tribe Papionini. <i>BMC Evolutionary Biology</i> , 2014, 14, 176.	3.2	49
41	The application of "omics" technologies for the classification and identification of animals. <i>Organisms Diversity and Evolution</i> , 2016, 16, 1-12.	1.6	49
42	The radiation of macaques out of Africa: Evidence from mitogenome divergence times and the fossil record. <i>Journal of Human Evolution</i> , 2019, 133, 114-132.	2.6	49
43	Phylogenetic Relationships among the Colobine Monkeys Revisited: New Insights from Analyses of Complete mt Genomes and 44 Nuclear Non-Coding Markers. <i>PLoS ONE</i> , 2012, 7, e36274.	2.5	48
44	Insights into the evolution of social systems and species from baboon studies. <i>ELife</i> , 2019, 8, .	6.0	47
45	The impact of storage buffer, DNA extraction method, and polymerase on microbial analysis. <i>Scientific Reports</i> , 2018, 8, 6292.	3.3	46
46	Phylogeny and distribution of crested gibbons (genus <i>Nomascus</i>) based on mitochondrial cytochrome b gene sequence data. <i>American Journal of Primatology</i> , 2010, 72, 1047-1054.	1.7	44
47	An Alu-Based Phylogeny of Gibbons (<i>Hylobatidae</i>). <i>Molecular Biology and Evolution</i> , 2012, 29, 3441-3450.	8.9	41
48	Nonhuman primates across sub-Saharan Africa are infected with the yaws bacterium <i>Treponema pallidum</i> subsp. <i>pertenue</i> . <i>Emerging Microbes and Infections</i> , 2018, 7, 1-4.	6.5	41
49	A novel landscape genetic approach demonstrates the effects of human disturbance on the Udzungwa red colobus monkey (<i>Procolobus gordonorum</i>). <i>Heredity</i> , 2016, 116, 167-176.	2.6	37
50	The Influence of Social Systems on Patterns of Mitochondrial DNA Variation in Baboons. <i>International Journal of Primatology</i> , 2014, 35, 210-225.	1.9	35
51	Genome sequence of the basal haplorrhine primate <i>Tarsius syrichta</i> reveals unusual insertions. <i>Nature Communications</i> , 2016, 7, 12997.	12.8	32
52	Widespread <i>Treponema pallidum</i> Infection in Nonhuman Primates, Tanzania. <i>Emerging Infectious Diseases</i> , 2018, 24, 1002-1009.	4.3	32
53	Inferring the evolutionary histories of divergences in <i>Hylobates</i> and <i>Nomascus</i> gibbons through multilocus sequence data. <i>BMC Evolutionary Biology</i> , 2013, 13, 82.	3.2	31
54	Population genetic insights into the social organization of Guinea baboons (<i>Papio papio</i>): Evidence for female-biased dispersal. <i>American Journal of Primatology</i> , 2015, 77, 878-889.	1.7	30

#	ARTICLE	IF	CITATIONS
55	A comparative analysis of Y chromosome and mtDNA phylogenies of the Hylobates gibbons. BMC Evolutionary Biology, 2012, 12, 150.	3.2	28
56	Population genetic structure of Guizhou snub-nosed monkeys (<i>Rhinopithecus brelichi</i>) as inferred from mitochondrial control region sequences, and comparison with <i>R. roxellana</i> and <i>R. bieti</i> . American Journal of Physical Anthropology, 2012, 147, 1-10.	2.1	28
57	Species-specific effects of climate change on the distribution of suitable baboon habitats – Ecological niche modeling of current and Last Glacial Maximum conditions. Journal of Human Evolution, 2019, 132, 215-226.	2.6	28
58	Isolation of Treponema DNA from Necrophagous Flies in a Natural Ecosystem. EBioMedicine, 2016, 11, 85-90.	6.1	27
59	An expanded mammal mitogenome dataset from Southeast Asia. GigaScience, 2017, 6, 1-8.	6.4	27
60	Population genomics of wild Chinese rhesus macaques reveals a dynamic demographic history and local adaptation, with implications for biomedical research. GigaScience, 2018, 7, .	6.4	27
61	Phylogeography, mitochondrial DNA diversity, and demographic history of geladas (Theropithecus) Tj ETQq1 1 0.784314 rgBT /Overlock 2.5 27	2.5	27
62	Genomic Mechanisms of Physiological and Morphological Adaptations of Limestone Langurs to Karst Habitats. Molecular Biology and Evolution, 2020, 37, 952-968.	8.9	27
63	Extrapair Paternity in Golden-Cheeked Gibbons (Nomascus gabriellae) in the Secondary Lowland Forest of Cat Tien National Park, Vietnam. Folia Primatologica, 2011, 82, 154-164.	0.7	26
64	Out of Africa, but how and when? The case of hamadryas baboons (Papio hamadryas). Journal of Human Evolution, 2014, 76, 154-164.	2.6	25
65	Introduction to Special Issue on Primate Hybridization and Hybrid Zones. International Journal of Primatology, 2019, 40, 1-8.	1.9	24
66	Inverted intergeneric introgression between critically endangered kipunjis and yellow baboons in two disjunct populations. Biology Letters, 2018, 14, 20170729.	2.3	23
67	Bonds of bros and brothers: Kinship and social bonding in postdispersal male macaques. Molecular Ecology, 2020, 29, 3346-3360.	3.9	23
68	Pan-African Voyagers: The Phylogeography of Baboons. , 2011, , 319-358.		22
69	Distribution of Mitochondrial Clades and Morphotypes of Baboons <i>Papio</i> spp. (Primates:) Tj ETQq1 1 0.784314 rgBT /Overlock 0.6 22	0.6	22
70	Implications of genetics and current protected areas for conservation of 5 endangered primates in China. Conservation Biology, 2015, 29, 1508-1517.	4.7	21
71	An Alu-Based Phylogeny of Lemurs (Infraorder: Lemuriformes). PLoS ONE, 2012, 7, e44035.	2.5	21
72	Considerable haplotypic diversity in the RT1-CE class I gene region of the rat major histocompatibility complex. Immunogenetics, 2005, 56, 773-777.	2.4	20

#	ARTICLE	IF	CITATIONS
73	Comparing chromosomal and mitochondrial phylogenies of sportive lemurs (Genus <i>Lepilemur</i> .) Tj ETQq1 1 0.784314.rgBT /Overlock 10	2.2	20
74	Comparing chromosomal and mitochondrial phylogenies of the Indriidae (Primates, Lemuriformes). <i>Chromosome Research</i> , 2011, 19, 209-224.	2.2	20
75	Genetic Diversity in Endangered Guizhou Snub-Nosed Monkeys (<i>Rhinopithecus brelichi</i>): Contrasting Results from Microsatellite and Mitochondrial DNA Data. <i>PLoS ONE</i> , 2013, 8, e73647.	2.5	20
76	Full-length <i>Numt</i> analysis provides evidence for hybridization between the Asian colobine genera <i>Trachypithecus</i> and <i>Semnopithecus</i> . <i>American Journal of Primatology</i> , 2015, 77, 901-910.	1.7	20
77	Highly polymorphic colour vision in a New World monkey with red facial skin, the bald uakari (<i>Cacajao calvus</i>). <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2016, 283, 20160067.	2.6	20
78	The Origin and Population History of the Endangered Golden Snub-Nosed Monkey (<i>Rhinopithecus</i>) Tj ETQq0 0 0.rgBT /Overlock 10 TF	8.9	20
79	Phylogenetic relationships of Malaysia's long-tailed macaques, <i>Macaca fascicularis</i> , based on cytochrome <i>b</i> sequences. <i>ZooKeys</i> , 2014, 407, 121-139.	1.1	19
80	Diversity and Evolutionary History of Macaques with Special Focus on <i>Macaca mulatta</i> and <i>Macaca fascicularis</i> . , 2015, , 3-16.		19
81	Population genetic structure and evolutionary history of Bale monkeys (<i>Chlorocebus djamdjamensis</i>) in the southern Ethiopian Highlands. <i>BMC Evolutionary Biology</i> , 2018, 18, 106.	3.2	18
82	Reconstructing the phylogeny of new world monkeys (platyrrhini): evidence from multiple non-coding loci. <i>Environmental Epigenetics</i> , 2019, 65, 579-588.	1.8	18
83	Metabarcoding of eukaryotic parasite communities describes diverse parasite assemblages spanning the primate phylogeny. <i>Molecular Ecology Resources</i> , 2020, 20, 204-215.	4.8	18
84	Relatively Recent Evolution of Pelage Coloration in Colobinae: Phylogeny and Phylogeography of Three Closely Related Langur Species. <i>PLoS ONE</i> , 2013, 8, e61659.	2.5	17
85	Insights into the genetic foundation of aggression in <i>Papio</i> and the evolution of two length-polymorphisms in the promoter regions of serotonin-related genes (5-HTTLPR and MAOALPR) in <i>Papionini</i> . <i>BMC Evolutionary Biology</i> , 2016, 16, 121.	3.2	17
86	The Tenasserim Lutung, <i>Trachypithecus barbei</i> (Blyth, 1847) (Primates: Cercopithecidae): Description of a live specimen, and a reassessment of phylogenetic affinities, taxonomic history, and distribution. <i>Contributions To Zoology</i> , 2004, 73, 271-282.	0.5	15
87	Discordance Between Spatial Distributions of Y-Chromosomal and Mitochondrial Haplotypes in African Green Monkeys (<i>Chlorocebus</i> spp.): A Result of Introgressive Hybridization or Cryptic Diversity?. <i>International Journal of Primatology</i> , 2013, 34, 986-999.	1.9	15
88	Differentiated adaptive evolution, episodic relaxation of selective constraints, and pseudogenization of umami and sweet taste genes TAS1Rs in catarrhine primates. <i>Frontiers in Zoology</i> , 2014, 11, 79.	2.0	15
89	Comparing mitogenomic timetrees for two African savannah primate genera (<i>Chlorocebus</i> and <i>Papio</i>). <i>Zoological Journal of the Linnean Society</i> , 2017, 181, 471-483.	2.3	15
90	Female Assamese macaques bias their affiliation to paternal and maternal kin. <i>Behavioral Ecology</i> , 2020, 31, 493-507.	2.2	15

#	ARTICLE	IF	CITATIONS
91	Population Genomics Reveals Incipient Speciation, Introgression, and Adaptation in the African Mona Monkey (<i>Cercopithecus mona</i>). <i>Molecular Biology and Evolution</i> , 2021, 38, 876-890.	8.9	15
92	Is <i>Colobus guereza gallarum</i> a valid endemic Ethiopian taxon?. <i>Primate Biology</i> , 2019, 6, 7-16.	1.0	15
93	Phylogeny and Classification of Gibbons (Hylobatidae). <i>Developments in Primatology</i> , 2016, , 151-165.	0.1	14
94	Strain diversity of <i>Treponema pallidum</i> subsp. <i>pertenue</i> suggests rare interspecies transmission in African nonhuman primates. <i>Scientific Reports</i> , 2019, 9, 14243.	3.3	14
95	Genetic Diversity, Inbreeding Level, and Genetic Load in Endangered Snub-Nosed Monkeys (<i>Rhinopithecus</i>). <i>Frontiers in Genetics</i> , 2020, 11, 615926.	2.3	14
96	Deep divergence among mitochondrial lineages in African jackals. <i>Zoologica Scripta</i> , 2018, 47, 1-8.	1.7	13
97	Genomic skimming and nanopore sequencing uncover cryptic hybridization in one of world's most threatened primates. <i>Scientific Reports</i> , 2021, 11, 17279.	3.3	13
98	Mitogenomic phylogeny of the Asian colobine genus <i>Trachypithecus</i> with special focus on <i>Trachypithecus phayrei</i> (Blyth, 1847) and description of a new species. <i>Zoological Research</i> , 2020, 41, 656-669.	2.1	13
99	A PCR-based marker to simply identify <i>Saimiri sciureus</i> and <i>S. boliviensis boliviensis</i> . <i>American Journal of Primatology</i> , 2008, 70, 1177-1180.	1.7	12
100	The Hybrid Origin of the Indochinese Gray Langur <i>Trachypithecus crepusculus</i> . <i>International Journal of Primatology</i> , 2019, 40, 9-27.	1.9	12
101	Genetic monogamy and mate choice in a pair-living primate. <i>Scientific Reports</i> , 2020, 10, 20328.	3.3	12
102	Reproductive parameters in Guizhou snub-nosed monkeys (<i>Rhinopithecus brelichi</i>). <i>American Journal of Primatology</i> , 2009, 71, 266-270.	1.7	11
103	So what is a species anyway? A primatological perspective. <i>Evolutionary Anthropology</i> , 2014, 23, 21-23.	3.4	11
104	<i>COMT</i> Val ¹⁵⁸ Met moderates the link between rank and aggression in a non-human primate. <i>Genes, Brain and Behavior</i> , 2018, 17, e12443.	2.2	11
105	The gut microbiome of exudivorous marmosets in the wild and captivity. <i>Scientific Reports</i> , 2022, 12, 5049.	3.3	11
106	The phylogenetic position of "Papio ruhei" a unique baboon taxon from Somalia. <i>Der Zoologische Garten</i> , 2008, 77, 303-311.	0.3	10
107	New mitogenomic lineages in <i>Papio</i> baboons and their phylogeographic implications. <i>American Journal of Physical Anthropology</i> , 2021, 174, 407-417.	2.1	10
108	Lactation and menstruation shift the vaginal microbiota in captive rhesus monkeys to be more similar to the male urethral microbiota. <i>Scientific Reports</i> , 2019, 9, 17399.	3.3	9

#	ARTICLE	IF	CITATIONS
109	Human Activity and Forest Degradation Threaten Populations of the Nigeriaâ€“Cameroon Chimpanzee (<i>Pan troglodytes ellioti</i>) in Western Cameroon. <i>International Journal of Primatology</i> , 2021, 42, 105-129.	1.9	8
110	Mitogenomic phylogeny of <i>Callithrix</i> with special focus on human transferred taxa. <i>BMC Genomics</i> , 2021, 22, 239.	2.8	8
111	Complete mitochondrial genome of a Toque Macaque (<i>Macaca sinica</i>). <i>Mitochondrial DNA Part B: Resources</i> , 2018, 3, 182-183.	0.4	7
112	Variation in predicted COVIDâ€“19 risk among lemurs and lorises. <i>American Journal of Primatology</i> , 2021, 83, e23255.	1.7	7
113	Initiation of the Primate Genome Project. <i>Zoological Research</i> , 2022, 43, 147-149.	2.1	7
114	Molecular phylogeny and systematics of bald uakaris, genus <i>Cacajao</i> (Primates: Pitheciidae), with the description of a new species. <i>Molecular Phylogenetics and Evolution</i> , 2022, 173, 107509.	2.7	7
115	Global view on virus infection in non-human primates and implications for public health and wildlife conservation. <i>Zoological Research</i> , 2021, 42, 626-632.	2.1	6
116	Environmental and anthropogenic effects on the nesting patterns of Nigeriaâ€“Cameroon chimpanzees in Northâ€“West Cameroon. <i>American Journal of Primatology</i> , 2021, 83, e23312.	1.7	6
117	Ancient DNA of the pygmy marmoset type specimen <i>Cebuella pygmaea</i> (Spix, 1823) resolves a taxonomic conundrum. <i>Zoological Research</i> , 2021, 42, 761-771.	2.1	6
118	Molecular Advances in Lorisid Taxonomy and Phylogeny. , 2020, , 57-66.		5
119	Mitogenomes of historical type specimens unravel the taxonomy of sportive lemurs (<i>Lepilemur</i> spp.) in Northwest Madagascar. <i>Zoological Research</i> , 2021, 42, 428-432.	2.1	5
120	Geographic distribution of microsatellite alleles in geladas (Primates, Cercopithecidae): Evidence for three evolutionary units. <i>Zoologica Scripta</i> , 2020, 49, 659-667.	1.7	4
121	Molecular Genetics Supports a Potential Fifth Asian Pangolin Species (Mammalia, Pholidota, Manis). <i>Zoological Science</i> , 2020, 37, 538-543.	0.7	4
122	Recently Integrated Alu Elements in Capuchin Monkeys: A Resource for <i>Cebus/Sapajus</i> Genomics. <i>Genes</i> , 2022, 13, 572.	2.4	4
123	Frequent non-reciprocal exchange in microsatellite-containing-DNA-regions of vertebrates. <i>Journal of Zoological Systematics and Evolutionary Research</i> , 2009, 47, 15-20.	1.4	3
124	Primate Taxonomy and Conservation. <i>Developments in Primatology</i> , 2016, , 193-213.	0.1	3
125	Complete mitochondrial genome of an olive baboon (<i>Papio anubis</i>) from Gombe National Park, Tanzania. <i>Mitochondrial DNA Part B: Resources</i> , 2018, 3, 177-178.	0.4	3
126	A New World Monkey Resembles Human in Bitter Taste Receptor Evolution and Function via a Single Parallel Amino Acid Substitution. <i>Molecular Biology and Evolution</i> , 2021, 38, 5472-5479.	8.9	3

#	ARTICLE	IF	CITATIONS
127	Novel mtDNA haplotypes represented in the European captive population of the Endangered François's™ langur (<i>Trachypithecus francoisi</i>). <i>International Journal of Primatology</i> , 2022, 43, 533-537.	1.9	3
128	Comparative analysis of DNA extraction protocols for ancient soft tissue museum samples. <i>Zoological Research</i> , 2021, 42, 280-286.	2.1	2
129	Population history of chimpanzees introduced to Lake Victoria's™ Rubondo Island. <i>Primates</i> , 2021, 62, 253-265.	1.1	2
130	Comparing mitogenomic timetrees for two African savannah primate genera (<i>Chlorocebus</i> and <i>Papio</i>). <i>Zoological Journal of the Linnean Society</i> , 2020, 190, 1071-1073.	2.3	1
131	Genotyping of non-human primate models: perspectives and challenges for the implementation of the "three R's". <i>Primate Biology</i> , 2014, 1, 1-9.	1.0	1
132	A refined panel of 42 microsatellite loci to universally genotype catarrhine primates. <i>Ecology and Evolution</i> , 2021, 11, 498-505.	1.9	1
133	Swayne's hartebeest in Ethiopia: population estimate, genetic variability and competition with livestock. <i>Oryx</i> , 0, , 1-9.	1.0	1
134	Mito-phylogenetic relationship of the new subspecies of gentle monkey <i>Cercopithecus mitis manyaraensis</i> , Butynski & De Jong, 2020. <i>Primate Biology</i> , 2022, 9, 11-18.	1.0	1
135	Non-invasive genotyping with a massively parallel sequencing panel for the detection of SNPs in HPA-axis genes. <i>Scientific Reports</i> , 2018, 8, 15944.	3.3	0
136	Taxonomic Revision and Evolutionary Phylogeography of Dusky Langur (<i>Presbytis dussumieri</i>) in Peninsular Malaysia. <i>Zoological Studies</i> , 2020, 59, e64.	0.3	0