

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

76031

42
h-index

71088

80
g-index

144
all docs

144
docs citations

144
times ranked

8161
citing authors

#	ARTICLE	IF	CITATIONS
1	Initiation of the Primate Genome Project. <i>Zoological Research</i> , 2022, 43, 147-149.	0.9	7
2	Recently Integrated Alu Elements in Capuchin Monkeys: A Resource for <i>Cebus/Sapajus</i> Genomics. <i>Genes</i> , 2022, 13, 572.	1.0	4
3	The gut microbiome of exudivorous marmosets in the wild and captivity. <i>Scientific Reports</i> , 2022, 12, 5049.	1.6	11
4	Novel mtDNA haplotypes represented in the European captive population of the Endangered FranÃ§oisâ€™ langur (<i>Trachypithecus francoisi</i>). <i>International Journal of Primatology</i> , 2022, 43, 533-537.	0.9	3
5	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.	1.2	7
6	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.	0.6	1
7	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.	3.5	15
8	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
9	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.	0.9	5
10	Comparative analysis of DNA extraction protocols for ancient soft tissue museum samples. <i>Zoological Research</i> , 2021, 42, 280-286.	0.9	2
11	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.	0.9	6
12	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.	0.9	8
13	Population history of chimpanzees introduced to Lake Victoriaâ€™s Rubondo Island. <i>Primates</i> , 2021, 62, 253-265.	0.7	2
14	Mitogenomic phylogeny of <i>Callithrix</i> with special focus on human transferred taxa. <i>BMC Genomics</i> , 2021, 22, 239.	1.2	8
15	Variation in predicted COVIDâ€™19 risk among lemurs and lorises. <i>American Journal of Primatology</i> , 2021, 83, e23255.	0.8	7
16	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.	0.8	6
17	Genomic skimming and nanopore sequencing uncover cryptic hybridization in one of worldâ€™s most threatened primates. <i>Scientific Reports</i> , 2021, 11, 17279.	1.6	13
18	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.	3.5	3

#	ARTICLE	IF	CITATIONS
19	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.	0.9	6
20	A refined panel of 42 microsatellite loci to universally genotype catarrhine primates. <i>Ecology and Evolution</i> , 2021, 11, 498-505.	0.8	1
21	Metabarcoding of eukaryotic parasite communities describes diverse parasite assemblages spanning the primate phylogeny. <i>Molecular Ecology Resources</i> , 2020, 20, 204-215.	2.2	18
22	Female Assamese macaques bias their affiliation to paternal and maternal kin. <i>Behavioral Ecology</i> , 2020, 31, 493-507.	1.0	15
23	Genomic Mechanisms of Physiological and Morphological Adaptations of Limestone Langurs to Karst Habitats. <i>Molecular Biology and Evolution</i> , 2020, 37, 952-968.	3.5	27
24	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.	1.0	1
25	Genetic monogamy and mate choice in a pair-living primate. <i>Scientific Reports</i> , 2020, 10, 20328.	1.6	12
26	Bonds of bros and brothers: Kinship and social bonding in postdispersal male macaques. <i>Molecular Ecology</i> , 2020, 29, 3346-3360.	2.0	23
27	Geographic distribution of microsatellite alleles in geladas (Primates, Cercopithecidae): Evidence for three evolutionary units. <i>Zoologica Scripta</i> , 2020, 49, 659-667.	0.7	4
28	A Severe Lack of Evidence Limits Effective Conservation of the World's Primates. <i>BioScience</i> , 2020, 70, 794-803.	2.2	51
29	Molecular Advances in Lorisid Taxonomy and Phylogeny. , 2020, , 57-66.		5
30	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.	0.9	13
31	Genetic Diversity, Inbreeding Level, and Genetic Load in Endangered Snub-Nosed Monkeys (<i>Rhinopithecus</i>). <i>Frontiers in Genetics</i> , 2020, 11, 615926.	1.1	14
32	Molecular Genetics Supports a Potential Fifth Asian Pangolin Species (Mammalia, Pholidota, Manis). <i>Zoological Science</i> , 2020, 37, 538-543.	0.3	4
33	Taxonomic Revision and Evolutionary Phylogeography of Dusky Langur (<i>Presbytis</i>) in Peninsular Malaysia. <i>Zoological Studies</i> , 2020, 59, e64.	0.3	0
34	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.	1.3	49
35	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.	1.6	14
36	The comparative genomics and complex population history of <i>Papio</i> baboons. <i>Science Advances</i> , 2019, 5, eaau6947.	4.7	115

#	ARTICLE	IF	CITATIONS
37	Hybridization in human evolution: Insights from other organisms. <i>Evolutionary Anthropology</i> , 2019, 28, 189-209.	1.7	57
38	Species-specific effects of climate change on the distribution of suitable baboon habitats – Ecological niche modeling of current and Last Glacial Maximum conditions. <i>Journal of Human Evolution</i> , 2019, 132, 215-226.	1.3	28
39	Introduction to Special Issue on Primate Hybridization and Hybrid Zones. <i>International Journal of Primatology</i> , 2019, 40, 1-8.	0.9	24
40	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.	1.6	9
41	The Origin and Population History of the Endangered Golden Snub-Nosed Monkey (<i>Rhinopithecus</i>) Tj ETQq1 1 0,784314 rgBT /Overl	3.5	20
42	Reconstructing the phylogeny of new world monkeys (platyrrhini): evidence from multiple non-coding loci. <i>Environmental Epigenetics</i> , 2019, 65, 579-588.	0.9	18
43	The Hybrid Origin of the Indochinese Gray Langur <i>Trachypithecus crepusculus</i> . <i>International Journal of Primatology</i> , 2019, 40, 9-27.	0.9	12
44	Is <i>Colobus guereza gallarum</i> a valid endemic Ethiopian taxon?. <i>Primate Biology</i> , 2019, 6, 7-16.	0.6	15
45	Insights into the evolution of social systems and species from baboon studies. <i>ELife</i> , 2019, 8, .	2.8	47
46	The impact of storage buffer, DNA extraction method, and polymerase on microbial analysis. <i>Scientific Reports</i> , 2018, 8, 6292.	1.6	46
47	Inverted intergeneric introgression between critically endangered kipunjis and yellow baboons in two disjunct populations. <i>Biology Letters</i> , 2018, 14, 20170729.	1.0	23
48	Complete mitochondrial genome of a Toque Macaque (<i>Macaca sinica</i>). <i>Mitochondrial DNA Part B: Resources</i> , 2018, 3, 182-183.	0.2	7
49	Deep divergence among mitochondrial lineages in African jackals. <i>Zoologica Scripta</i> , 2018, 47, 1-8.	0.7	13
50	<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.	1.1	11
51	Primates in peril: the significance of Brazil, Madagascar, Indonesia and the Democratic Republic of the Congo for global primate conservation. <i>PeerJ</i> , 2018, 6, e4869.	0.9	123
52	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.	3.0	41
53	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.	1.6	0
54	Population genomics of wild Chinese rhesus macaques reveals a dynamic demographic history and local adaptation, with implications for biomedical research. <i>GigaScience</i> , 2018, 7, .	3.3	27

#	ARTICLE	IF	CITATIONS
55	Widespread <i>Treponema pallidum</i> Infection in Nonhuman Primates, Tanzania. <i>Emerging Infectious Diseases</i> , 2018, 24, 1002-1009.	2.0	32
56	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
57	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.	0.2	3
58	Phylogeography, mitochondrial DNA diversity, and demographic history of geladas (<i>Theropithecus</i>)	1.1	27
59	Impending extinction crisis of the world's primates: Why primates matter. <i>Science Advances</i> , 2017, 3, e1600946.	4.7	912
60	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.	0.8	80
61	Species definitions and conservation: a review and case studies from African mammals. <i>Conservation Genetics</i> , 2017, 18, 1247-1256.	0.8	58
62	Morphometric, Behavioral, and Genomic Evidence for a New Orangutan Species. <i>Current Biology</i> , 2017, 27, 3487-3498.e10.	1.8	192
63	An expanded mammal mitogenome dataset from Southeast Asia. <i>GigaScience</i> , 2017, 6, 1-8.	3.3	27
64	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.	1.0	15
65	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 Papionini. <i>BMC Evolutionary Biology</i> , 2016, 16, 121.	3.2	17
66	Taxonomic review of the New World tamarins (Primates: Callitrichidae). <i>Zoological Journal of the Linnean Society</i> , 2016, 177, 1003-1028.	1.0	59
67	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.	1.2	20
68	Isolation of <i>Treponema</i> DNA from Necrophagous Flies in a Natural Ecosystem. <i>EBioMedicine</i> , 2016, 11, 85-90.	2.7	27
69	Primate Taxonomy and Conservation. <i>Developments in Primatology</i> , 2016, , 193-213.	0.7	3
70	Phylogeny and Classification of Gibbons (Hylobatidae). <i>Developments in Primatology</i> , 2016, , 151-165.	0.7	14
71	Sustained virologic control in SIV macaques after antiretroviral and antibody therapy. <i>Science</i> , 2016, 354, 197-202.	6.0	194
72	Genome sequence of the basal haplorrhine primate <i>Tarsius syrichta</i> reveals unusual insertions. <i>Nature Communications</i> , 2016, 7, 12997.	5.8	32

#	ARTICLE	IF	CITATIONS
73	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.	1.2	37
74	The application of "omics" technologies for the classification and identification of animals. <i>Organisms Diversity and Evolution</i> , 2016, 16, 1-12.	0.7	49
75	Remarkable ancient divergences amongst neglected loriform primates. <i>Zoological Journal of the Linnean Society</i> , 2015, 175, 661-674.	1.0	71
76	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.	0.8	30
77	Distribution of Mitochondrial Clades and Morphotypes of Baboons <i>Papio</i> spp. (Primates: Cercopithecoidea). <i>Molecular Biology and Evolution</i> , 2015, 32, 1000-1010.	0.6	22
78	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.	0.8	20
79	Mitogenomic phylogeny of the common long-tailed macaque (<i>Macaca fascicularis fascicularis</i>). <i>BMC Genomics</i> , 2015, 16, 222.	1.2	55
80	Implications of genetics and current protected areas for conservation of 5 endangered primates in China. <i>Conservation Biology</i> , 2015, 29, 1508-1517.	2.4	21
81	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.	1.2	50
82	Diversity and Evolutionary History of Macaques with Special Focus on <i>Macaca mulatta</i> and <i>Macaca fascicularis</i> . <i>Journal of Biogeography</i> , 2015, 42, 3-16.		19
83	Mitogenomics of the Old World monkey tribe Papionini. <i>BMC Evolutionary Biology</i> , 2014, 14, 176.	3.2	49
84	Out of Africa, but how and when? The case of hamadryas baboons (<i>Papio hamadryas</i>). <i>Journal of Human Evolution</i> , 2014, 76, 154-164.	1.3	25
85	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.	0.5	19
86	Genome typing of nonhuman primate models: implications for biomedical research. <i>Trends in Genetics</i> , 2014, 30, 482-487.	2.9	54
87	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.	0.9	15
88	The Influence of Social Systems on Patterns of Mitochondrial DNA Variation in Baboons. <i>International Journal of Primatology</i> , 2014, 35, 210-225.	0.9	35
89	Whole-genome sequencing of the snub-nosed monkey provides insights into folivory and evolutionary history. <i>Nature Genetics</i> , 2014, 46, 1303-1310.	9.4	174
90	So what is a species anyway? A primatological perspective. <i>Evolutionary Anthropology</i> , 2014, 23, 21-23.	1.7	11

#	ARTICLE	IF	CITATIONS
91	Gibbon genome and the fast karyotype evolution of small apes. <i>Nature</i> , 2014, 513, 195-201.	13.7	320
92	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.	0.6	1
93	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
94	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.	0.9	15
95	Mitochondrial Diversity and Distribution of African Green Monkeys (<i>Chlorocebus</i> Gray, 1870). <i>American Journal of Primatology</i> , 2013, 75, 350-360.	0.8	87
96	Baboon phylogeny as inferred from complete mitochondrial genomes. <i>American Journal of Physical Anthropology</i> , 2013, 150, 133-140.	2.1	110
97	Relatively Recent Evolution of Pelage Coloration in Colobinae: Phylogeny and Phylogeography of Three Closely Related Langur Species. <i>PLoS ONE</i> , 2013, 8, e61659.	1.1	17
98	A Mitogenomic Phylogeny of Living Primates. <i>PLoS ONE</i> , 2013, 8, e69504.	1.1	217
99	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.	1.1	20
100	An Alu-Based Phylogeny of Gibbons (<i>Hylobatidae</i>). <i>Molecular Biology and Evolution</i> , 2012, 29, 3441-3450.	3.5	41
101	A comparative analysis of Y chromosome and mtDNA phylogenies of the <i>Hylobates</i> gibbons. <i>BMC Evolutionary Biology</i> , 2012, 12, 150.	3.2	28
102	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.	1.1	48
103	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.	1.1	53
104	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> . <i>American Journal of Physical Anthropology</i> , 2012, 147, 1-10.	2.1	28
105	An Alu-Based Phylogeny of Lemurs (Infraorder: Lemuriformes). <i>PLoS ONE</i> , 2012, 7, e44035.	1.1	21
106	Extrapair Paternity in Golden-Cheeked Gibbons (<i>Nomascus gabriellae</i>) in the Secondary Lowland Forest of Cat Tien National Park, Vietnam. <i>Folia Primatologica</i> , 2011, 82, 154-164.	0.3	26
107	Pan-African Voyagers: The Phylogeography of Baboons. , 2011, , 319-358.		22
108	A Molecular Phylogeny of Living Primates. <i>PLoS Genetics</i> , 2011, 7, e1001342.	1.5	1,130

#	ARTICLE	IF	CITATIONS
109	Nuclear versus mitochondrial DNA: evidence for hybridization in colobine monkeys. BMC Evolutionary Biology, 2011, 11, 77.	3.2	123
110	Comparing chromosomal and mitochondrial phylogenies of the Indriidae (Primates, Lemuriformes). Chromosome Research, 2011, 19, 209-224.	1.0	20
111	Concordance between vocal and genetic diversity in crested gibbons. BMC Evolutionary Biology, 2011, 11, 36.	3.2	75
112	Mitochondrial phylogeny of tamarins (<i>Saguinus</i> , Hoffmannsegg 1807) with taxonomic and biogeographic implications for the <i>S. nigricollis</i> species group. American Journal of Physical Anthropology, 2011, 144, 564-574.	2.1	65
113	The strange blood: Natural hybridization in primates. Evolutionary Anthropology, 2011, 20, 96-103.	1.7	146
114	Mitochondrial phylogeny of leaf monkeys (genus <i>Presbytis</i> , Eschscholtz, 1821) with implications for taxonomy and conservation. Molecular Phylogenetics and Evolution, 2011, 59, 311-319.	1.2	49
115	Introgressive hybridization in southern African baboons shapes patterns of mtDNA variation. American Journal of Physical Anthropology, 2010, 142, 125-136.	2.1	52
116	Mitochondrial evidence for multiple radiations in the evolutionary history of small apes. BMC Evolutionary Biology, 2010, 10, 74.	3.2	111
117	Phylogeny and distribution of crested gibbons (genus <i>Nomascus</i>) based on mitochondrial cytochrome b gene sequence data. American Journal of Primatology, 2010, 72, 1047-1054.	0.8	44
118	Mitochondrial Genome Sequences Effectively Reveal the Phylogeny of Hylobates Gibbons. PLoS ONE, 2010, 5, e14419.	1.1	75
119	Is the New Primate Genus <i>Rungwecebus</i> a Baboon?. PLoS ONE, 2009, 4, e4859.	1.1	66
120	Mitochondrial phylogeography of baboons (<i>Papio</i> spp.) – Indication for introgressive hybridization?. BMC Evolutionary Biology, 2009, 9, 83.	3.2	173
121	Reproductive parameters in Guizhou snub-nosed monkeys (<i>Rhinopithecus brelichi</i>). American Journal of Primatology, 2009, 71, 266-270.	0.8	11
122	Frequent non-reciprocal exchange in microsatellite-containing-DNA-regions of vertebrates. Journal of Zoological Systematics and Evolutionary Research, 2009, 47, 15-20.	0.6	3
123	Retropositional events consolidate the branching order among New World monkey genera. Molecular Phylogenetics and Evolution, 2009, 50, 507-513.	1.2	60
124	Comparing chromosomal and mitochondrial phylogenies of sportive lemurs (Genus <i>Lepilemur</i>). Journal of Zoological Systematics and Evolutionary Research, 2009, 47, 15-20.	1.0	20
125	A PCR-based marker to simply identify <i>Saimiri sciureus</i> and <i>S. boliviensis boliviensis</i> . American Journal of Primatology, 2008, 70, 1177-1180.	0.8	12
126	Mitochondrial phylogeny, taxonomy and biogeography of the silvered langur species group (<i>Trachypithecus cristatus</i>). Molecular Phylogenetics and Evolution, 2008, 47, 629-636.	1.2	53

#	ARTICLE	IF	CITATIONS
127	The phylogenetic position of "Papio ruhei" a unique baboon taxon from Somalia. Der Zoologische Garten, 2008, 77, 303-311.	0.3	10
128	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
129	Molecular phylogeny and evolutionary history of Southeast Asian macaques forming the M. silenus group. Molecular Phylogenetics and Evolution, 2007, 42, 807-816.	1.2	89
130	Molecular phylogeny and taxonomic revision of the sportive lemurs (Lepilemur, Primates). BMC Evolutionary Biology, 2006, 6, 17.	3.2	59
131	Considerable haplotypic diversity in the RT1-CE class I gene region of the rat major histocompatibility complex. Immunogenetics, 2005, 56, 773-777.	1.2	20
132	Primate phylogeny: molecular evidence from retroposons. Cytogenetic and Genome Research, 2005, 108, 26-37.	0.6	66
133	The Tenasserim Lutung, Trachypithecus barbei (Blyth, 1847) (Primates: Cercopithecidae): Description of a live specimen, and a reassessment of phylogenetic affinities, taxonomic history, and distribution. Contributions To Zoology, 2004, 73, 271-282.	0.2	15
134	Primate jumping genes elucidate strepsirrhine phylogeny. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 10650-10654.	3.3	238
135	Molecular Phylogeny of the Major Hylobatid Divisions. Molecular Phylogenetics and Evolution, 2001, 19, 486-494.	1.2	103
136	Swayne's hartebeest in Ethiopia: population estimate, genetic variability and competition with livestock. Oryx, 0, , 1-9.	0.5	1