

# Christian Roos

## List of Publications by Year in descending order

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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	Initiation of the Primate Genome Project. <i>Zoological Research</i> , 2022, 43, 147-149.	2.1	7
2	Recently Integrated Alu Elements in Capuchin Monkeys: A Resource for <i>Cebus/Sapajus</i> Genomics. <i>Genes</i> , 2022, 13, 572.	2.4	4
3	The gut microbiome of exudivorous marmosets in the wild and captivity. <i>Scientific Reports</i> , 2022, 12, 5049.	3.3	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.	1.9	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.	2.7	7
6	Mito-phylogenetic relationship of the new subspecies of gentle monkey &lt;i>Cercopithecus mitis manyaraensis&lt;/i>, Butynski & De Jong, 2020. <i>Primate Biology</i> , 2022, 9, 11-18.	1.0	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.	8.9	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 (&lt;i>Lepilemur&lt;/i> spp.) in Northwest Madagascar. <i>Zoological Research</i> , 2021, 42, 428-432.	2.1	5
10	Comparative analysis of DNA extraction protocols for ancient soft tissue museum samples. <i>Zoological Research</i> , 2021, 42, 280-286.	2.1	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.	2.1	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.	1.9	8
13	Population history of chimpanzees introduced to Lake Victoriaâ€™s Rubondo Island. <i>Primates</i> , 2021, 62, 253-265.	1.1	2
14	Mitogenomic phylogeny of <i>Callithrix</i> with special focus on human transferred taxa. <i>BMC Genomics</i> , 2021, 22, 239.	2.8	8
15	Variation in predicted COVIDâ€™19 risk among lemurs and lorises. <i>American Journal of Primatology</i> , 2021, 83, e23255.	1.7	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.	1.7	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.	3.3	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.	8.9	3

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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.	2.1	6
20	A refined panel of 42 microsatellite loci to universally genotype catarrhine primates. <i>Ecology and Evolution</i> , 2021, 11, 498-505.	1.9	1
21	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
22	Female Assamese macaques bias their affiliation to paternal and maternal kin. <i>Behavioral Ecology</i> , 2020, 31, 493-507.	2.2	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.	8.9	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.	2.3	1
25	Genetic monogamy and mate choice in a pair-living primate. <i>Scientific Reports</i> , 2020, 10, 20328.	3.3	12
26	Bonds of bros and brothers: Kinship and social bonding in postdispersal male macaques. <i>Molecular Ecology</i> , 2020, 29, 3346-3360.	3.9	23
27	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
28	A Severe Lack of Evidence Limits Effective Conservation of the World's Primates. <i>BioScience</i> , 2020, 70, 794-803.	4.9	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.	2.1	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.	2.3	14
32	Molecular Genetics Supports a Potential Fifth Asian Pangolin Species (Mammalia, Pholidota, Manis). <i>Zoological Science</i> , 2020, 37, 538-543.	0.7	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.	2.6	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.	3.3	14
36	The comparative genomics and complex population history of <i>Papio</i> baboons. <i>Science Advances</i> , 2019, 5, eaau6947.	10.3	115

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37	Hybridization in human evolution: Insights from other organisms. <i>Evolutionary Anthropology</i> , 2019, 28, 189-209.	3.4	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.	2.6	28
39	Introduction to Special Issue on Primate Hybridization and Hybrid Zones. <i>International Journal of Primatology</i> , 2019, 40, 1-8.	1.9	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.	3.3	9
41	The Origin and Population History of the Endangered Golden Snub-Nosed Monkey ( <i>Rhinopithecus</i> )	8.9	20
42	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
43	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
44	Is <i>Colobus guereza gallarum</i> a valid endemic Ethiopian taxon?. <i>Primate Biology</i> , 2019, 6, 7-16.	1.0	15
45	Insights into the evolution of social systems and species from baboon studies. <i>ELife</i> , 2019, 8, .	6.0	47
46	The impact of storage buffer, DNA extraction method, and polymerase on microbial analysis. <i>Scientific Reports</i> , 2018, 8, 6292.	3.3	46
47	Inverted intergeneric introgression between critically endangered kipunjis and yellow baboons in two disjunct populations. <i>Biology Letters</i> , 2018, 14, 20170729.	2.3	23
48	Complete mitochondrial genome of a Toque Macaque ( <i>Macaca sinica</i> ). <i>Mitochondrial DNA Part B: Resources</i> , 2018, 3, 182-183.	0.4	7
49	Deep divergence among mitochondrial lineages in African jackals. <i>Zoologica Scripta</i> , 2018, 47, 1-8.	1.7	13
50	<i>COMT</i> Val <sup>158</sup> 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
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.	2.0	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.	6.5	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.	3.3	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, .	6.4	27

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55	Widespread <i>Treponema pallidum</i> Infection in Nonhuman Primates, Tanzania. <i>Emerging Infectious Diseases</i> , 2018, 24, 1002-1009.	4.3	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.	0.4	3
58	Phylogeography, mitochondrial DNA diversity, and demographic history of geladas ( <i>Theropithecus</i> ) Tj ETQq0 0 0 rgBT/Overlock 10 Tf 50	2.5	27
59	Impending extinction crisis of the world's primates: Why primates matter. <i>Science Advances</i> , 2017, 3, e1600946.	10.3	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.	1.7	80
61	Species definitions and conservation: a review and case studies from African mammals. <i>Conservation Genetics</i> , 2017, 18, 1247-1256.	1.5	58
62	Morphometric, Behavioral, and Genomic Evidence for a New Orangutan Species. <i>Current Biology</i> , 2017, 27, 3487-3498.e10.	3.9	192
63	An expanded mammal mitogenome dataset from Southeast Asia. <i>GigaScience</i> , 2017, 6, 1-8.	6.4	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.	2.3	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.	2.3	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.	2.6	20
68	Isolation of <i>Treponema</i> DNA from Necrophagous Flies in a Natural Ecosystem. <i>EBioMedicine</i> , 2016, 11, 85-90.	6.1	27
69	Primate Taxonomy and Conservation. <i>Developments in Primatology</i> , 2016, , 193-213.	0.1	3
70	Phylogeny and Classification of Gibbons (Hylobatidae). <i>Developments in Primatology</i> , 2016, , 151-165.	0.1	14
71	Sustained virologic control in SIV <sup>+</sup> macaques after antiretroviral and $\hat{\pm}$ antibody therapy. <i>Science</i> , 2016, 354, 197-202.	12.6	194
72	Genome sequence of the basal haplorrhine primate <i>Tarsius syrichta</i> reveals unusual insertions. <i>Nature Communications</i> , 2016, 7, 12997.	12.8	32

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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.	2.6	37
74	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
75	Remarkable ancient divergences amongst neglected loriform primates. <i>Zoological Journal of the Linnean Society</i> , 2015, 175, 661-674.	2.3	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.	1.7	30
77	Distribution of Mitochondrial Clades and Morphotypes of Baboons <i>Papio</i> spp. (Primates). <i>Trends in Ecology and Evolution</i> , 2015, 30, 100-106.	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.	1.7	20
79	Mitogenomic phylogeny of the common long-tailed macaque ( <i>Macaca fascicularis fascicularis</i> ). <i>BMC Genomics</i> , 2015, 16, 222.	2.8	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.	4.7	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.	2.7	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.	2.6	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.	1.1	19
86	Genome typing of nonhuman primate models: implications for biomedical research. <i>Trends in Genetics</i> , 2014, 30, 482-487.	6.7	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.	2.0	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.	1.9	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.	21.4	174
90	So what is a species anyway? A primatological perspective. <i>Evolutionary Anthropology</i> , 2014, 23, 21-23.	3.4	11

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91	Gibbon genome and the fast karyotype evolution of small apes. <i>Nature</i> , 2014, 513, 195-201.	27.8	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.	1.0	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.	1.9	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.	1.7	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.	2.5	17
98	A Mitogenomic Phylogeny of Living Primates. <i>PLoS ONE</i> , 2013, 8, e69504.	2.5	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.	2.5	20
100	An Alu-Based Phylogeny of Gibbons ( <i>Hylobatidae</i> ). <i>Molecular Biology and Evolution</i> , 2012, 29, 3441-3450.	8.9	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.	2.5	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.	2.5	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 (Infraclass: Lemuriformes). <i>PLoS ONE</i> , 2012, 7, e44035.	2.5	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.	0.7	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.	3.5	1,130

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109	Nuclear versus mitochondrial DNA: evidence for hybridization in colobine monkeys. <i>BMC Evolutionary Biology</i> , 2011, 11, 77.	3.2	123
110	Comparing chromosomal and mitochondrial phylogenies of the Indriidae (Primates, Lemuriformes). <i>Chromosome Research</i> , 2011, 19, 209-224.	2.2	20
111	Concordance between vocal and genetic diversity in crested gibbons. <i>BMC Evolutionary Biology</i> , 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. <i>American Journal of Physical Anthropology</i> , 2011, 144, 564-574.	2.1	65
113	The strange blood: Natural hybridization in primates. <i>Evolutionary Anthropology</i> , 2011, 20, 96-103.	3.4	146
114	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
115	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
116	Mitochondrial evidence for multiple radiations in the evolutionary history of small apes. <i>BMC Evolutionary Biology</i> , 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. <i>American Journal of Primatology</i> , 2010, 72, 1047-1054.	1.7	44
118	Mitochondrial Genome Sequences Effectively Reveal the Phylogeny of <i>Hylobates</i> Gibbons. <i>PLoS ONE</i> , 2010, 5, e14419.	2.5	75
119	Is the New Primate Genus <i>Rungwecebus</i> a Baboon?. <i>PLoS ONE</i> , 2009, 4, e4859.	2.5	66
120	Mitochondrial phylogeography of baboons ( <i>Papio</i> spp.) – Indication for introgressive hybridization?. <i>BMC Evolutionary Biology</i> , 2009, 9, 83.	3.2	173
121	Reproductive parameters in Guizhou snub-nosed monkeys ( <i>Rhinopithecus brelichi</i> ). <i>American Journal of Primatology</i> , 2009, 71, 266-270.	1.7	11
122	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
123	Retropositional events consolidate the branching order among New World monkey genera. <i>Molecular Phylogenetics and Evolution</i> , 2009, 50, 507-513.	2.7	60
124	Comparing chromosomal and mitochondrial phylogenies of sportive lemurs (Genus <i>Lepilemur</i> ). <i>Journal of Molecular Evolution</i> , 2009, 69, 10-14.	2.2	20
125	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
126	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



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127	The phylogenetic position of "Papio ruhei" a unique baboon taxon from Somalia. <i>Der Zoologische Garten</i> , 2008, 77, 303-311.	0.3	10
128	Phylogenetic position of the langur genera <i>Semnopithecus</i> and <i>Trachypithecus</i> among Asian colobines, and genus affiliations of their species groups. <i>BMC Evolutionary Biology</i> , 2008, 8, 58.	3.2	94
129	Molecular phylogeny and evolutionary history of Southeast Asian macaques forming the <i>M. silenus</i> group. <i>Molecular Phylogenetics and Evolution</i> , 2007, 42, 807-816.	2.7	89
130	Molecular phylogeny and taxonomic revision of the sportive lemurs ( <i>Lepilemur</i> , Primates). <i>BMC Evolutionary Biology</i> , 2006, 6, 17.	3.2	59
131	Considerable haplotypic diversity in the RT1-CE class I gene region of the rat major histocompatibility complex. <i>Immunogenetics</i> , 2005, 56, 773-777.	2.4	20
132	Primate phylogeny: molecular evidence from retroposons. <i>Cytogenetic and Genome Research</i> , 2005, 108, 26-37.	1.1	66
133	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
134	Primate jumping genes elucidate strepsirrhine phylogeny. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004, 101, 10650-10654.	7.1	238
135	Molecular Phylogeny of the Major Hylobatid Divisions. <i>Molecular Phylogenetics and Evolution</i> , 2001, 19, 486-494.	2.7	103
136	Swayne's hartebeest in Ethiopia: population estimate, genetic variability and competition with livestock. <i>Oryx</i> , 0, , 1-9.	1.0	1