Sandro L Bonatto

List of Publications by Year in descending order

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130

all docs

129 5,951 39 papers citations h-index

130

docs citations

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130 6525
times ranked citing authors

71

#	Article	IF	CITATIONS
1	So close, so far: spatial genetic structure and mating system in <i>Petunia exserta</i> , an endemic from a peculiar landscape in the Brazilian Pampa grasslands. Botanical Journal of the Linnean Society, 2022, 199, 412-427.	1.6	7
2	Genetic diversity in micro-endemic plants from highland grasslands in southern Brazil. Botanical Journal of the Linnean Society, 2022, 199, 235-251.	1.6	6
3	Gone With the Water: The Loss of Genetic Variability in Black and Gold Howler Monkeys (Alouatta) Tj ETQq1 1 ().784314 2.2	rgBT /Overlock
4	Population Genetics and Phylogeography of Galapagos Fur Seals. Frontiers in Genetics, 2022, 13, .	2.3	1
5	When phylogeography meets niche suitability to unravel the evolutionary history of a shrub from the Brazilian Atlantic Forest. Botanical Journal of the Linnean Society, 2021, 195, 77-92.	1.6	3
6	Phylogenomic Discordance in the Eared Seals is best explained by Incomplete Lineage Sorting following Explosive Radiation in the Southern Hemisphere. Systematic Biology, 2021, 70, 786-802.	5.6	25
7	How diverse can rare species be on the margins of genera distribution?. AoB PLANTS, 2019, 11, plz037.	2.3	12
8	Phylogeographic evidence for two species of muriqui (genus <i>Brachyteles</i>). American Journal of Primatology, 2019, 81, e23066.	1.7	21
9	Southern extension of the geographic range of black-and-gold howler monkeys (<i>Alouatta) Tj ETQq1 1 0.784.</i>	314 rgBT /	Overlock 10 Tf
10	Contact zones and their consequences: hybridization between two ecologically isolated wild Petunia species. Botanical Journal of the Linnean Society, 2019, , .	1.6	1
11	Population structure, phylogeography, and genetic diversity of the common bottlenose dolphin in the tropical and subtropical southwestern Atlantic Ocean. Journal of Mammalogy, 2019, 100, 564-577.	1.3	17
12	Molecular phylogeny and hemipenial diversity of South American species of <i>Amerotyphlops </i> (Typhlopidae, Scolecophidia). Zoologica Scripta, 2019, 48, 139-156.	1.7	13
13	Hybridization Between Neotropical Primates with Contrasting Sexual Dichromatism. International Journal of Primatology, 2019, 40, 99-113.	1.9	14
14	Temporal stability and mixed-stock analyses of humpback whales (Megaptera novaeangliae) in the nearshore waters of the Western Antarctic Peninsula. Polar Biology, 2018, 41, 323-340.	1.2	14
15	How strong was the bottleneck associated to the peopling of the Americas? New insights from multilocus sequence data. Genetics and Molecular Biology, 2018, 41, 206-214.	1.3	31
16	Effective population size and the genetic consequences of commercial whaling on the humpback whales (Megaptera novaeangliae) from Southwestern Atlantic Ocean. Genetics and Molecular Biology, 2018, 41, 253-262.	1.3	3
17	Origin and hidden diversity within the poorly known Gal $ ilde{A}_i$ pagos snake radiation (Serpentes:) Tj ETQq $1\ 1\ 0.7845$	314 rgBT / 1.2	Overlock 10 Tf
18	Phylogeny and systematics of Chiroxiphia and Antilophia manakins (Aves, Pipridae). Molecular Phylogenetics and Evolution, 2018, 127, 706-711.	2.7	12

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19	From inland to the coast: Spatial and environmental signatures on the genetic diversity in the colonization of the South Atlantic Coastal Plain. Perspectives in Plant Ecology, Evolution and Systematics, 2017, 28, 47-57.	2.7	18
20	Rare or cryptic? The first report of an Omura's whale (<i>Balaenoptera omurai</i>) in the South Atlantic Ocean. Marine Mammal Science, 2017, 33, 80-95.	1.8	14
21	Genetic differentiation between humpback whales (<i>Megaptera novaeangliae</i>) from Atlantic and Pacific breeding grounds of South America. Marine Mammal Science, 2017, 33, 457-479.	1.8	13
22	Secondary structure of nrDNA Internal Transcribed Spacers as a useful tool to align highly divergent species in phylogenetic studies. Genetics and Molecular Biology, 2017, 40, 191-199.	1.3	7
23	Ancient female philopatry, asymmetric male gene flow, and synchronous population expansion support the influence of climatic oscillations on the evolution of South American sea lion (Otaria) Tj $\rm ETQq1~1~0.$	7842 3 4 rg	gBT ⊉@verlock
24	Multiple introductions and gene flow in subtropical South American populations of the fireweed, Senecio madagascariensis(Asteraceae). Genetics and Molecular Biology, 2016, 39, 135-144.	1.3	14
25	Effects of past climate on (i) Passiflora actinia (i) (Passifloraceae) populations and insights into future species management in the Brazilian Atlantic forest. Botanical Journal of the Linnean Society, 2016, 180, 348-364.	1.6	11
26	Multiple evolutionary units and demographic stability during the last glacial maximum in the Scytalopus speluncae complex (Aves: Rhinocryptidae). Molecular Phylogenetics and Evolution, 2016, 102, 86-96.	2.7	15
27	Could refuge theory and rivers acting as barriers explain the genetic variability distribution in the Atlantic Forest?. Molecular Phylogenetics and Evolution, 2016, 101, 242-251.	2.7	49
28	High levels of genetic diversity and population structure in an endemic and rare species: implications for conservation. AoB PLANTS, 2016, 8, .	2.3	52
29	Discovery of a chemosynthesis-based community in the western South Atlantic Ocean. Deep-Sea Research Part I: Oceanographic Research Papers, 2016, 112, 45-56.	1.4	34
30	Ancient remains and the first peopling of the <scp>A</scp> mericas: Reassessing the Hoyo Negro skull. American Journal of Physical Anthropology, 2015, 158, 514-521.	2.1	28
31	Genetic differentiation and hybrid identification using microsatellite markers in closely related wild species. AoB PLANTS, 2015, 7, plv084.	2.3	47
32	Testing the effect of palaeodrainages versus habitat stability on genetic divergence in riverine systems: study of a Neotropical fish of the Brazilian coastal Atlantic Forest. Journal of Biogeography, 2015, 42, 2389-2401.	3.0	90
33	Re-evaluation of the generic status of <i> Athenaea < /i > and <i> Aureliana < /i > (Withaniinae, Solanaceae) based on molecular phylogeny and morphology of the calyx. Botanical Journal of the Linnean Society, 2015, 177, 322-334.</i></i>	1.6	12
34	Pollen dispersal and breeding structure in a hawkmoth-pollinated Pampa grasslands speciesPetunia axillaris(Solanaceae). Annals of Botany, 2015, 115, 939-948.	2.9	37
35	Fine-scale matrilineal population structure in the Galapagos fur seal and its implications for conservation management. Conservation Genetics, 2015, 16, 1099-1113.	1.5	25
36	Were sea level changes during the Pleistocene in the South Atlantic Coastal Plain a driver of speciation in Petunia (Solanaceae)?. BMC Evolutionary Biology, 2015, 15, 92.	3.2	33

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37	Novel Microsatellites for Calibrachoa heterophylla (Solanaceae) Endemic to the South Atlantic Coastal Plain of South America. Applications in Plant Sciences, 2015, 3, 1500021.	2.1	5
38	Novel Transposable Elements in Solanaceae: Evolutionary Relationships among Tnt1-related Sequences in Wild Petunia Species. Plant Molecular Biology Reporter, 2014, 32, 142-152.	1.8	11
39	Phylogeography of the <i>Petunia integrifolia </i> complex in southern Brazil. Botanical Journal of the Linnean Society, 2014, 174, 199-213.	1.6	34
40	Molecular insights into the purpleâ€flowered ancestor of garden petunias. American Journal of Botany, 2014, 101, 119-127.	1.7	24
41	Nuclear and plastid markers reveal the persistence of genetic identity: A new perspective on the evolutionary history of Petunia exserta. Molecular Phylogenetics and Evolution, 2014, 70, 504-512.	2.7	42
42	Genetic diversity and ecological niche modelling of the restrictedRecordia reitzii(Verbenaceae) from southern Brazilian Atlantic forest. Botanical Journal of the Linnean Society, 2014, 176, 332-348.	1.6	22
43	Multilocus phylogeny reconstruction: New insights into the evolutionary history of the genus Petunia. Molecular Phylogenetics and Evolution, 2014, 81, 19-28.	2.7	63
44	Diversification in the <scp>S</scp> outh <scp>A</scp> merican <scp>P</scp> ampas: the genetic and morphological variation of the widespread <i><scp>P</scp>etunia axillaris</i> complex (<scp>S</scp> olanaceae). Molecular Ecology, 2014, 23, 374-389.	3.9	54
45	Reconciling pre-Columbian settlement hypotheses requires integrative, multidisciplinary, and model-bound approaches. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, E213-4.	7.1	18
46	Geological and climatic changes in quaternary shaped the evolutionary history of Calibrachoa heterophylla, an endemic South-Atlantic species of petunia. BMC Evolutionary Biology, 2013, 13, 178.	3.2	35
47	Development of Microsatellites forVerbenoxylum reitzii(Verbenaceae), a Tree Endemic to the Brazilian Atlantic Forest. Applications in Plant Sciences, 2013, 1, 1300005.	2.1	1
48	Molecular systematics and historical biogeography of tree boas (Corallus spp.). Molecular Phylogenetics and Evolution, 2013, 66, 953-959.	2.7	36
49	Biogeographical history and diversification of <i>Petunia </i> li>and <i>Calibrachoa </i> (Solanaceae) in the Neotropical Pampas grassland. Botanical Journal of the Linnean Society, 2013, 171, 140-153.	1.6	77
50	A Molecular Systematic Analysis of Passiflora ovalis and Passiflora contracta (Passifloraceae). Phytotaxa, 2013, 132, 39.	0.3	8
51	Influence of the 48867A>C (Asp358Ala) IL6R polymorphism on response to a lifestyle modification intervention in individuals with metabolic syndrome. Genetics and Molecular Research, 2013, 12, 3983-3991.	0.2	11
52	A Bayesian Approach to Genome/Linguistic Relationships in Native South Americans. PLoS ONE, 2013, 8, e64099.	2.5	12
53	Isolation and Characterization of Microsatellite Markers for Passiflora contracta. International Journal of Molecular Sciences, 2012, 13, 11343-11348.	4.1	16
54	Cultural diversification promotes rapid phenotypic evolution in Xavante Indians. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 73-77.	7.1	17

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55	Phylogeny, biogeography and divergence times in Passiflora (Passifloraceae). Genetics and Molecular Biology, 2012, 35, 1036-1043.	1.3	59
56	Microsatellites in Aureliana fasciculatavar. fasciculata (Solanaceae), a shrub that inhabits the Atlantic Rainforest. American Journal of Botany, 2012, 99, e173-e175.	1.7	0
57	An Alternative Model for the Early Peopling of Southern South America Revealed by Analyses of Three Mitochondrial DNA Haplogroups. PLoS ONE, 2012, 7, e43486.	2.5	88
58	Infrageneric classification of <i>Calibrachoa</i> (Solanaceae) based on morphological and molecular evidence. Taxon, 2012, 61, 120-130.	0.7	22
59	Molecular phylogeny of the New World Dipsadidae (Serpentes: Colubroidea): a reappraisal. Cladistics, 2012, 28, 437-459.	3.3	112
60	Conservation genetics of South American aquatic mammals: an overview of gene diversity, population structure, phylogeography, nonâ€invasive methods and forensics. Mammal Review, 2012, 42, 275-303.	4.8	10
61	The population genetics of quechuas, the largest native south american group: Autosomal sequences, SNPs, and microsatellites evidence high level of diversity. American Journal of Physical Anthropology, 2012, 147, 443-451.	2.1	11
62	Isolation, characterization, and crossâ€amplification of microsatellite markers for the <i>Petunia integrifolia</i> (Solanaceae) complex. American Journal of Botany, 2011, 98, e277-9.	1.7	10
63	Permanent Genetic Resources added to Molecular Ecology Resources Database 1 June 2011–31 July 2011. Molecular Ecology Resources, 2011, 11, 1124-1126.	4.8	14
64	Distribution of Yâ€chromosome q lineages in native americans. American Journal of Human Biology, 2011, 23, 563-566.	1.6	26
65	A new subhaplogroup of native American Y-Chromosomes from the Andes. American Journal of Physical Anthropology, 2011, 146, 553-559.	2.1	38
66	Molecular phylogeny of the Neoplecostominae and Hypoptopomatinae (Siluriformes: Loricariidae) using multiple genes. Molecular Phylogenetics and Evolution, 2011, 59, 43-52.	2.7	77
67	High rate of viral evolution in the capsid protein of porcine parvovirus. Journal of General Virology, 2011, 92, 2628-2636.	2.9	52
68	Does Variation in Genome Sizes Reflect Adaptive or Neutral Processes? New Clues from Passiflora. PLoS ONE, 2011, 6, e18212.	2.5	52
69	The use and limits of ITS data in the analysis of intraspecific variation in Passiflora L. (Passifloraceae). Genetics and Molecular Biology, 2010, 33, 99-108.	1.3	36
70	The phylogenetic placement of Hollandichthys Eigenmann 1909 (Teleostei: Characidae) and related genera. Molecular Phylogenetics and Evolution, 2010, 57, 1347-1352.	2.7	18
71	Diversification of plant species in a subtropical region of eastern South American highlands: a phylogeographic perspective on native <i>Petunia</i> (Solanaceae). Molecular Ecology, 2010, 19, 5240-5251.	3.9	75

Molecular phylogeny of the South American land slug < i> Phyllocaulis < /i> (Mollusca, Soleolifera,) Tj ETQq $0\ 0\ 0\ rgBT$ / Oyerlock 10 Tf 50 62

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73	Mitochondrial control region haplotypes of the South American sea lion Otaria flavescens (Shaw,) Tj ETQq $1\ 1\ 0.7$	784314 rgl	3T/Overlock 14
74	A functional ABCA1 gene variant is associated with low HDL-cholesterol levels and shows evidence of positive selection in Native Americans. Human Molecular Genetics, 2010, 19, 2877-2885.	2.9	133
7 5	Microsatellite Genetic Characterization of the Humpback Whale (Megaptera novaeangliae) Breeding Ground off Brazil (Breeding Stock A). Journal of Heredity, 2010, 101, 189-200.	2.4	24
76	Molecular phylogeny of advanced snakes (Serpentes, Caenophidia) with an emphasis on South American Xenodontines: a revised classification and descriptions of new taxa. Papeis Avulsos De Zoologia, 2009, 49, 115-153.	0.4	262
77	Population Structure of Humpback Whales from Their Breeding Grounds in the South Atlantic and Indian Oceans. PLoS ONE, 2009, 4, e7318.	2.5	84
78	Characterization of new microsatellite loci for the South-American rodents Cavia aperea and C. magna. Conservation Genetics Resources, 2009, 1, 47-50.	0.8	9
79	Multigene phylogeny and DNA barcoding indicate that the Sandwich tern complex (Thalasseus) Tj ETQq1 1 0.78 52, 263-267.	4314 rgBT 2.7	Overlock 10 20
80	Molecular phylogeny and biogeography of the eastern Tapaculos (Aves: Rhinocryptidae: Scytalopus,) Tj ETQq0 0 Evolution, 2009, 53, 450-462.	0 rgBT /Ο\ 2.7	verlock 10 Tf 46
81	Population data of 17 Y-STR loci from Rio Grande do Sul state (South Brazil). Forensic Science International: Genetics, 2009, 4, e31-e33.	3.1	14
82	Isolation and characterization of 12 dinucletiotide microsatellite loci in <i>Paratrechalea galianoae</i> (Araneae, Trechaleidae), a nuptial giftâ€spider. Molecular Ecology Resources, 2009, 9, 539-541.	4.8	4
83	First molecular estimate of sex-ratio of southern right whale calves, Eubalaena australis, for Brazilian waters. Journal of the Marine Biological Association of the United Kingdom, 2009, 89, 1003-1007.	0.8	5
84	Uniparental (mtDNA, Yâ€chromosome) Polymorphisms in French Guiana and Two Related Populations – Implications for the Region's Colonization. Annals of Human Genetics, 2008, 72, 145-156.	0.8	24
85	Mitochondrial DNA diversity of the Southwestern Atlantic humpback whale (Megaptera novaeangliae) breeding area off Brazil, and the potential connections to Antarctic feeding areas. Conservation Genetics, 2008, 9, 1253-1262.	1.5	38
86	The peopling of America: Craniofacial shape variation on a continental scale and its interpretation from an interdisciplinary view. American Journal of Physical Anthropology, 2008, 137, 175-187.	2.1	163
87	Hidden generic diversity in Neotropical birds: Molecular and anatomical data support a new genus for the "Scytalopus―indigoticus species-group (Aves: Rhinocryptidae). Molecular Phylogenetics and Evolution, 2008, 49, 125-135.	2.7	25
88	Mitochondrial Population Genomics Supports a Single Pre-Clovis Origin with a Coastal Route for the Peopling of the Americas. American Journal of Human Genetics, 2008, 82, 583-592.	6.2	319
89	Reply to Ho and Endicott. American Journal of Human Genetics, 2008, 83, 146-147.	6.2	19
90	DNA sequence analysis and the phylogeographical history of the rodent <i>Deltamys kempi</i> (Sigmodontinae, Cricetidae) on the Atlantic Coastal Plain of south of Brazil. Journal of Evolutionary Biology, 2008, 21, 1823-1835.	1.7	9

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91	Interâ€species hybridization among Neotropical cats of the genus <i>Leopardus</i> , and evidence for an introgressive hybrid zone between <i>L. geoffroyi</i> and <i>L. tigrinus</i> in southern Brazil. Molecular Ecology, 2008, 17, 4317-4333.	3.9	83
92	Mapping the evolutionary twilight zone: molecular markers, populations and geography. Journal of Biogeography, 2008, 35, 753-763.	3.0	61
93	Crossâ€amplification and characterization of 13 tetranucleotide microsatellites in multiple species of Neotropical canids. Molecular Ecology Resources, 2008, 8, 898-900.	4.8	8
94	Taxonomy of Ixinandria Isbrýcker & Nijssen (Loricariidae: Loricariinae) based on morphological and molecular data. Neotropical Ichthyology, 2008, 6, 367-378.	1.0	5
95	A Reevaluation of the Native American MtDNA Genome Diversity and Its Bearing on the Models of Early Colonization of Beringia. PLoS ONE, 2008, 3, e3157.	2.5	60
96	Statistical evaluation of alternative models of human evolution. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 17614-17619.	7.1	497
97	Analysis of nucleotide diversity of NAT2 coding region reveals homogeneity across Native American populations and high intra-population diversity. Pharmacogenomics Journal, 2007, 7, 144-152.	2.0	42
98	Phylogenetic position of Placozoa based on large subunit (LSU) and small subunit (SSU) rRNA genes. Genetics and Molecular Biology, 2007, 30, 127-132.	1.3	40
99	Molecular Variability of the 16p13.3 Region in Amerindians and its Anthropological Significance. Annals of Human Genetics, 2007, 71, 64-76.	0.8	9
100	A phylogenomic appraisal of the evolutionary relationship of mycoplasmas. Genetics and Molecular Biology, 2007, 30, 270-276.	1.3	1
101	Aluinsertion polymorphisms in Native Americans and related Asian populations. Annals of Human Biology, 2006, 33, 142-160.	1.0	31
102	Molecular genetic variation in Passiflora alata (Passifloraceae), an invasive species in southern Brazil. Biological Journal of the Linnean Society, 2006, 88, 611-630.	1.6	12
103	Phylogeography of theBothrops jararacacomplex (Serpentes: Viperidae): past fragmentation and island colonization in the Brazilian Atlantic Forest. Molecular Ecology, 2006, 15, 3969-3982.	3.9	183
104	Diversity and natural hybridization in a highly endemic species of Petunia (Solanaceae): a molecular and ecological analysis. Molecular Ecology, 2006, 15, 4487-4497.	3.9	86
105	Molecular Modeling of Pathogenesis-Related Proteins of Family 5. Cell Biochemistry and Biophysics, 2006, 44, 385-394.	1.8	18
106	Molecular Phylogenetic Analysis of Petunia Juss. (Solanaceae). Genetica, 2006, 126, 3-14.	1.1	61
107	Differential organellar inheritance in Passiflora's (Passifloraceae) subgenera. Genetica, 2006, 128, 449-453.	1.1	17
108	Phylogenetic information in polymorphic L1 andAlu insertions from East Asians and Native American populations. American Journal of Physical Anthropology, 2005, 128, 171-184.	2.1	18

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109	Is haplogroup X present in extant South American Indians?. American Journal of Physical Anthropology, 2005, 127, 439-448.	2.1	34
110	Worldwide Genetic Variation at the 3′â€UTR Region of the <i>LDLR</i> Gene: Possible Influence of Natural Selection. Annals of Human Genetics, 2005, 69, 389-400.	0.8	17
111	Patterns of molecular evolution in pathogenesis-related proteins. Genetics and Molecular Biology, 2005, 28, 645-653.	1.3	20
112	Human T-cell lymphotropic virus type II in GuaranÃ-Indians, Southern Brazil. Cadernos De Saude Publica, 2005, 21, 1947-1951.	1.0	14
113	Swine and Poultry Pathogens: the Complete Genome Sequences of Two Strains of <i>Mycoplasma hyopneumoniae </i> and a Strain of <i>Mycoplasma synoviae </i> . Journal of Bacteriology, 2005, 187, 5568-5577.	2.2	289
114	Phylogeographic Inferences Concerning Evolution of Brazilian Passiflora actinia and P. elegans (Passifloraceae) Based on ITS (nrDNA) Variation. Annals of Botany, 2005, 95, 799-806.	2.9	52
115	B-FDNA sequence variability in Brazilian (blue-egg Caipira) chickens. Animal Genetics, 2004, 35, 278-284.	1.7	22
116	Molecular phylogeny of Trichomonadidae family inferred from ITS-1, 5.8S rRNA and ITS-2 sequences. International Journal for Parasitology, 2004, 34, 963-970.	3.1	73
117	Mitochondrial DNA and <i>Alu</i> insertions in a genetically peculiar population: The Ayoreo Indians of Bolivia and Paraguay. American Journal of Human Biology, 2004, 16, 479-488.	1.6	40
118	Short ReportExtremely limited mitochondrial DNA variability among the Aché Natives of Paraguay. Annals of Human Biology, 2004, 31, 87-94.	1.0	33
119	A first molecular phylogenetic analysis of <i>Passiflora </i> (Passifloraceae). American Journal of Botany, 2003, 90, 1229-1238.	1.7	99
120	Correction: Mitochondrial DNA Variation in Amerindians. American Journal of Human Genetics, 2003, 72, 1346-1348.	6.2	19
121	The complete genome sequence of Chromobacterium violaceum reveals remarkable and exploitable bacterial adaptability. Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 11660-11665.	7.1	251
122	Alu insertions versus blood group plus protein genetic variability in four Amerindian populations. Annals of Human Biology, 2002, 29, 334-347.	1.0	31
123	Mitochondrial Genome Diversity of Native Americans Supports a Single Early Entry of Founder Populations into America. American Journal of Human Genetics, 2002, 71, 187-192.	6.2	93
124	Genetic, geographic, and linguistic variation among South American Indians: Possible sex influence. American Journal of Physical Anthropology, 2002, 117, 68-78.	2.1	34
125	Extreme homogeneity among Brazilian wheat genotypes determined by RAPD markers. Pesquisa Agropecuaria Brasileira, 2000, 35, 2255-2260.	0.9	11
126	Phylogeographic Patterns and Evolution of the Mitochondrial DNA Control Region in Two Neotropical Cats (Mammalia, Felidae). Journal of Molecular Evolution, 1998, 47, 613-624.	1.8	87

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127	Diversity and Age of the Four Major mtDNA Haplogroups, and Their Implications for the Peopling of the New World. American Journal of Human Genetics, 1997, 61, 1413-1423.	6.2	128
128	A single and early migration for the peopling of the Americas supported by mitochondrial DNA sequence data. Proceedings of the National Academy of Sciences of the United States of America, 1997, 94, 1866-1871.	7.1	215
129	Identification and inter-relationship analysis of Bradyrhizobium japonicum strains by restriction fragment length polymorphism (RFLP) and random amplified polymorphic DNA (RAPD). World Journal of Microbiology and Biotechnology, 1994, 10, 648-652.	3.6	7