

Michael W. Bruford

List of Publications by Citations

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332 papers	18,616 citations	69 h-index	127 g-index
356 ext. papers	21,770 ext. citations	6.3 avg, IF	6.28 L-index

#	Paper	IF	Citations
332	Whole-genome analyses resolve early branches in the tree of life of modern birds. <i>Science</i> , 2014 , 346, 1320-31	33.3	1182
331	The sequence and de novo assembly of the giant panda genome. <i>Nature</i> , 2010 , 463, 311-7	50.4	864
330	Ecology. Essential biodiversity variables. <i>Science</i> , 2013 , 339, 277-8	33.3	809
329	Comparative genomics reveals insights into avian genome evolution and adaptation. <i>Science</i> , 2014 , 346, 1311-20	33.3	628
328	A Role for Ecotones in Generating Rainforest Biodiversity. <i>Science</i> , 1997 , 276, 1855-1857	33.3	509
327	DNA fingerprinting in birds. <i>Nature</i> , 1987 , 327, 149-52	50.4	485
326	Parental care and mating behaviour of polyandrous dunnocks <i>Prunella modularis</i> related to paternity by DNA fingerprinting. <i>Nature</i> , 1989 , 338, 249-251	50.4	448
325	Microsatellites and their application to population genetic studies. <i>Current Opinion in Genetics and Development</i> , 1993 , 3, 939-43	4.9	371
324	A spatial analysis method (SAM) to detect candidate loci for selection: towards a landscape genomics approach to adaptation. <i>Molecular Ecology</i> , 2007 , 16, 3955-69	5.7	345
323	DNA markers reveal the complexity of livestock domestication. <i>Nature Reviews Genetics</i> , 2003 , 4, 900-1030.1	30.1	345
322	Genomics and the challenging translation into conservation practice. <i>Trends in Ecology and Evolution</i> , 2015 , 30, 78-87	10.9	335
321	Revealing the history of sheep domestication using retrovirus integrations. <i>Science</i> , 2009 , 324, 532-6	33.3	292
320	Behavior predicts genes structure in a wild primate group. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1996 , 93, 5797-801	11.5	285
319	Genetic diversity and introgression in the Scottish wildcat. <i>Molecular Ecology</i> , 2001 , 10, 319-36	5.7	268
318	Genetic signature of anthropogenic population collapse in orang-utans. <i>PLoS Biology</i> , 2006 , 4, e25	9.7	212
317	Immigration and the ephemerality of a natural population bottleneck: evidence from molecular markers. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2001 , 268, 1387-94	4.4	211
316	SNeP: a tool to estimate trends in recent effective population size trajectories using genome-wide SNP data. <i>Frontiers in Genetics</i> , 2015 , 6, 109	4.5	184

315	Ecological constraints drive social evolution in the African mole-rats. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 1997 , 264, 1619-27	4.4	184
314	Rapid screening of invertebrate predators for multiple prey DNA targets. <i>Molecular Ecology</i> , 2005 , 14, 819-27	5.7	179
313	DNA answers the call of pipistrelle bat species. <i>Nature</i> , 1997 , 387, 138-9	50.4	174
312	Genetic divergence and units for conservation in the Komodo dragon <i>Varanus komodoensis</i> . <i>Proceedings of the Royal Society B: Biological Sciences</i> , 1999 , 266, 2269-2274	4.4	167
311	'Ghost' alleles of the Mauritius kestrel. <i>Nature</i> , 2000 , 403, 616	50.4	166
310	Genetic analysis reveals the wild ancestors of the llama and the alpaca. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2001 , 268, 2575-84	4.4	161
309	Molecular censusing doubles giant panda population estimate in a key nature reserve. <i>Current Biology</i> , 2006 , 16, R451-2	6.3	158
308	Patterns of genetic diversity and migration in increasingly fragmented and declining orang-utan (<i>Pongo pygmaeus</i>) populations from Sabah, Malaysia. <i>Molecular Ecology</i> , 2005 , 14, 441-56	5.7	152
307	Male bimaturism and reproductive success in Sumatran orang-utans. <i>Behavioral Ecology</i> , 2002 , 13, 643-652		152
306	The role of Pleistocene refugia and rivers in shaping gorilla genetic diversity in central Africa. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007 , 104, 20432-6	11.5	151
305	Large-scale mitochondrial DNA analysis of the domestic goat reveals six haplogroups with high diversity. <i>PLoS ONE</i> , 2007 , 2, e1012	3.7	145
304	Genetic diversity and subdivision of 57 European and Middle-Eastern sheep breeds. <i>Animal Genetics</i> , 2007 , 38, 37-44	2.5	133
303	Whole-genome sequencing of the snub-nosed monkey provides insights into folivory and evolutionary history. <i>Nature Genetics</i> , 2014 , 46, 1303-10	36.3	122
302	Permanent Genetic Resources added to Molecular Ecology Resources Database 1 May 2009-31 July 2009. <i>Molecular Ecology Resources</i> , 2009 , 9, 1460-6	8.4	122
301	Peregrine and saker falcon genome sequences provide insights into evolution of a predatory lifestyle. <i>Nature Genetics</i> , 2013 , 45, 563-6	36.3	119
300	A multi-samples, multi-extracts approach for microsatellite analysis of faecal samples in an arboreal ape. <i>Conservation Genetics</i> , 2000 , 1, 157-162	2.6	114
299	Convergent genomic signatures of domestication in sheep and goats. <i>Nature Communications</i> , 2018 , 9, 813	17.4	112
298	Reproductive skew among males in a female-dominated mammalian society. <i>Behavioral Ecology</i> , 2002 , 13, 193-200	2.3	112

297	Molecular technologies for biodiversity evaluation: opportunities and challenges. <i>Nature Biotechnology</i> , 1997 , 15, 625-8	44.5	108
296	Genetic structure of European sheep breeds. <i>Heredity</i> , 2007 , 99, 620-31	3.6	103
295	Genetic viability and population history of the giant panda, putting an end to the "evolutionary dead end"?. <i>Molecular Biology and Evolution</i> , 2007 , 24, 1801-10	8.3	100
294	Black and white and read all over: the past, present and future of giant panda genetics. <i>Molecular Ecology</i> , 2012 , 21, 5660-74	5.7	99
293	Philopatry and reproductive success in Bornean orang-utans (<i>Pongo pygmaeus</i>). <i>Molecular Ecology</i> , 2006 , 15, 2577-88	5.7	95
292	Sociogenetic structure in a free-living nocturnal primate population: sex-specific differences in the grey mouse lemur (<i>Microcebus murinus</i>). <i>Behavioral Ecology and Sociobiology</i> , 2001 , 50, 493-502	2.5	93
291	Estimation of admixture proportions: a likelihood-based approach using Markov chain Monte Carlo. <i>Genetics</i> , 2001 , 158, 1347-62	4	93
290	Set ambitious goals for biodiversity and sustainability. <i>Science</i> , 2020 , 370, 411-413	33.3	92
289	Bringing genetic diversity to the forefront of conservation policy and management. <i>Conservation Genetics Resources</i> , 2013 , 5, 593-598	0.8	91
288	Genetic diversity targets and indicators in the CBD post-2020 Global Biodiversity Framework must be improved. <i>Biological Conservation</i> , 2020 , 248, 108654	6.2	89
287	Turtle groups or turtle soup: dispersal patterns of hawksbill turtles in the Caribbean. <i>Molecular Ecology</i> , 2009 , 18, 4841-53	5.7	88
286	Microsatellite analysis of genetic diversity in fragmented South African buffalo populations. <i>Animal Conservation</i> , 1998 , 1, 85-94	3.2	88
285	Mitogenomic Meta-Analysis Identifies Two Phases of Migration in the History of Eastern Eurasian Sheep. <i>Molecular Biology and Evolution</i> , 2015 , 32, 2515-33	8.3	85
284	Riverine effects on mitochondrial structure of Bornean orangutans (<i>Pongo pygmaeus</i>) at two spatial scales. <i>Molecular Ecology</i> , 2008 , 17, 2898-909	5.7	84
283	Molecular tools and analytical approaches for the characterization of farm animal genetic diversity. <i>Animal Genetics</i> , 2012 , 43, 483-502	2.5	82
282	New perspectives on mate choice and the MHC. <i>Heredity</i> , 1998 , 81 (Pt 2), 127-33	3.6	82
281	Molecular detection of predation by soil micro-arthropods on nematodes. <i>Molecular Ecology</i> , 2006 , 15, 1963-72	5.7	82
280	The persistence of Pliocene populations through the Pleistocene climatic cycles: evidence from the phylogeography of an Iberian lizard. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2001 , 268, 1625-30	4.4	81

279	Molecular systematics and phylogeography of the cryptic species complex <i>Baetis rhodani</i> (Ephemeroptera, Baetidae). <i>Molecular Phylogenetics and Evolution</i> , 2006 , 40, 370-82	4.1	80
278	Parentage, reproductive skew and queen turnover in a multiple-queen ant analysed with microsatellites. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 1997 , 264, 277-83	4.4	79
277	The development of microsatellite loci in the song sparrow, <i>Melospiza melodia</i> (Aves) and genotyping errors associated with good quality DNA. <i>Molecular Ecology Notes</i> , 2001 , 1, 11-13		79
276	Permanent Genetic Resources added to Molecular Ecology Resources Database 1 October 2009-30 November 2009. <i>Molecular Ecology Resources</i> , 2010 , 10, 404-8	8.4	78
275	Molecular evidence for deep phylogenetic divergence in <i>Mandrillus sphinx</i> . <i>Molecular Ecology</i> , 2003 , 12, 2019-24	5.7	78
274	High performance computation of landscape genomic models including local indicators of spatial association. <i>Molecular Ecology Resources</i> , 2017 , 17, 1072-1089	8.4	77
273	Comparative evaluation of potential indicators and temporal sampling protocols for monitoring genetic erosion. <i>Evolutionary Applications</i> , 2014 , 7, 984-98	4.8	77
272	Primate conservation: measuring and mitigating trade in primates. <i>Endangered Species Research</i> , 2011 , 13, 159-161	2.5	75
271	The Value of Ecosystem Services from Giant Panda Reserves. <i>Current Biology</i> , 2018 , 28, 2174-2180.e7	6.3	75
270	Micro- and macrogeographical genetic structure of colonies of naked mole-rats <i>Heterocephalus glaber</i> . <i>Molecular Ecology</i> , 1997 , 6, 615-28	5.7	74
269	Gene-flow patterns in Atlantic and Mediterranean populations of the Lusitanian sea star <i>Asterina gibbosa</i> . <i>Molecular Ecology</i> , 2005 , 14, 3373-82	5.7	74
268	Projecting genetic diversity and population viability for the fragmented orang-utan population in the Kinabatangan floodplain, Sabah, Malaysia. <i>Endangered Species Research</i> , 2010 , 12, 249-261	2.5	74
267	Genetic and genomic monitoring with minimally invasive sampling methods. <i>Evolutionary Applications</i> , 2018 , 11, 1094-1119	4.8	72
266	The role of vicariance vs. dispersal in shaping genetic patterns in ocellated lizard species in the western Mediterranean. <i>Molecular Ecology</i> , 2008 , 17, 1535-51	5.7	72
265	Genetic evidence for female-biased dispersal and gene flow in a polygynous primate. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2006 , 273, 479-84	4.4	72
264	Unravelling migratory connectivity in marine turtles using multiple methods. <i>Journal of Applied Ecology</i> , 2010 , 47, 769-778	5.8	71
263	Molecular biogeography: towards an integrated framework for conserving pan-African biodiversity. <i>PLoS ONE</i> , 2007 , 2, e454	3.7	69
262	Polygynandry in a red fox population: implications for the evolution of group living in canids?. <i>Behavioral Ecology</i> , 2004 , 15, 766-778	2.3	69

261	Mating system and reproductive skew in the black rhinoceros. <i>Molecular Ecology</i> , 2001 , 10, 2031-41	5.7	69
260	Molecular analysis of dispersal in giant pandas. <i>Molecular Ecology</i> , 2007 , 16, 3792-800	5.7	67
259	Patterns and dynamics of sex-biased dispersal in a nocturnal primate, the grey mouse lemur, <i>Microcebus murinus</i> . <i>Animal Behaviour</i> , 2003 , 65, 709-719	2.8	65
258	Post-2020 goals overlook genetic diversity. <i>Science</i> , 2020 , 367, 1083-1085	33.3	64
257	Mitochondrial DNA phylogeography of western lowland gorillas (<i>Gorilla gorilla gorilla</i>). <i>Molecular Ecology</i> , 2004 , 13, 1551-65, 1567	5.7	64
256	Testing the reliability of microsatellite typing from faecal DNA in the savannah baboon. <i>Conservation Genetics</i> , 2000 , 1, 173-176	2.6	63
255	The Preservation of Process: The Missing Element of Conservation Programs. <i>Biodiversity Letters</i> , 1993 , 1, 164		63
254	Effective population size dynamics and the demographic collapse of Bornean orang-utans. <i>PLoS ONE</i> , 2012 , 7, e49429	3.7	62
253	Genetic structure of fragmented populations of red squirrel (<i>Sciurus vulgaris</i>) in the UK. <i>Molecular Ecology</i> , 1999 , 8, S55-63	5.7	62
252	Mitochondrial DNA diversity and phylogeography of endangered green turtle (<i>Chelonia mydas</i>) populations in Africa. <i>Conservation Genetics</i> , 2006 , 7, 353-369	2.6	61
251	Human microsatellites applicable for analysis of genetic variation in apes and Old World monkeys. <i>Journal of Heredity</i> , 1996 , 87, 406-10	2.4	61
250	Next-generation metrics for monitoring genetic erosion within populations of conservation concern. <i>Evolutionary Applications</i> , 2018 , 11, 1066-1083	4.8	59
249	Demographic loss, genetic structure and the conservation implications for Indian tigers. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2013 , 280, 20130496	4.4	59
248	DNA sequence variation and methylation in an arsenic tolerant earthworm population. <i>Soil Biology and Biochemistry</i> , 2013 , 57, 524-532	7.5	58
247	Species-specific mitochondrial DNA markers for identification of non-invasive samples from sympatric carnivores in the Iberian Peninsula. <i>Conservation Genetics</i> , 2008 , 9, 681-690	2.6	58
246	Biodiversity vs. biocontrol: positive and negative effects of alternative prey on control of slugs by carabid beetles. <i>Bulletin of Entomological Research</i> , 2006 , 96, 637-45	1.7	58
245	Ant workers selfishly bias sex ratios by manipulating female development. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2002 , 269, 173-8	4.4	57
244	Genetic structure and gene flow among Komodo dragon populations inferred by microsatellite loci analysis. <i>Molecular Ecology</i> , 1999 , 8, S17-30	5.7	57

243	Isolation of Microsatellite Markers in Animals 1998 , 279-285		56
242	Intraguild predation in winter wheat: prey choice by a common epigeal carabid consuming spiders. <i>Journal of Applied Ecology</i> , 2013 , 50, 271-279	5.8	55
241	New perspectives on mate choice and the MHC. <i>Heredity</i> , 1998 , 81 (Pt 3), 239-45	3.6	55
240	Sustaining genetic variation in a small population: evidence from the Mauritius kestrel. <i>Molecular Ecology</i> , 2001 , 10, 593-602	5.7	55
239	Genetic consequences of historical anthropogenic and ecological events on giant pandas. <i>Ecology</i> , 2013 , 94, 2346-57	4.6	54
238	Molecular evidence for Pleistocene refugia at the eastern edge of the Tibetan Plateau. <i>Molecular Ecology</i> , 2011 , 20, 3014-26	5.7	53
237	Crossing the Red Sea: phylogeography of the hamadryas baboon, <i>Papio hamadryas hamadryas</i> . <i>Molecular Ecology</i> , 2004 , 13, 2819-27	5.7	53
236	Molecular phylogeny and morphological change in the <i>Psittacula</i> parakeets. <i>Molecular Phylogenetics and Evolution</i> , 2004 , 31, 96-108	4.1	53
235	Permanent Genetic Resources added to Molecular Ecology Resources database 1 January 2009-30 April 2009. <i>Molecular Ecology Resources</i> , 2009 , 9, 1375-9	8.4	52
234	Prospects and challenges for the conservation of farm animal genomic resources, 2015-2025. <i>Frontiers in Genetics</i> , 2015 , 6, 314	4.5	50
233	Landscape, habitat characteristics and the genetic population structure of two caddisflies. <i>Freshwater Biology</i> , 2007 , 52, 1907-1929	3.1	50
232	Genomic signatures of adaptive introgression from European mouflon into domestic sheep. <i>Scientific Reports</i> , 2017 , 7, 7623	4.9	49
231	Whole-genome resequencing of wild and domestic sheep identifies genes associated with morphological and agronomic traits. <i>Nature Communications</i> , 2020 , 11, 2815	17.4	48
230	Modification of river meandering by tropical deforestation. <i>Geology</i> , 2017 , 45, 511-514	5	46
229	Landscape genomics and biased FST approaches reveal single nucleotide polymorphisms under selection in goat breeds of North-East Mediterranean. <i>BMC Genetics</i> , 2009 , 10, 7	2.6	46
228	Differential enzyme targeting as an evolutionary adaptation to herbivory in carnivora. <i>Molecular Biology and Evolution</i> , 2004 , 21, 632-46	8.3	46
227	Cross-species amplification, non-invasive genotyping, and non-Mendelian inheritance of human STRPs in Savannah baboons. <i>American Journal of Primatology</i> , 2000 , 51, 219-27	2.5	46
226	Odorant receptor gene expression changes during the parr-smolt transformation in Atlantic salmon. <i>Molecular Ecology</i> , 2004 , 13, 2851-7	5.7	45

225	Using nested clade analysis to assess the history of colonization and the persistence of populations of an Iberian Lizard. <i>Molecular Ecology</i> , 2002 , 11, 809-19	5.7	45
224	Phylogenetic Reanalysis of the Saudi Gazelle and Its Implications for Conservation. <i>Conservation Biology</i> , 2001 , 15, 1123-1133	6	45
223	Multilocus and single locus minisatellite analysis in population biological studies. <i>Exs</i> , 1991 , 58, 154-68		45
222	Inbreeding of bottlenecked butterfly populations. Estimation using the likelihood of changes in marker allele frequencies. <i>Genetics</i> , 1999 , 151, 1053-63	4	45
221	Conservation implications of drastic reductions in the smallest and most isolated populations of giant pandas. <i>Conservation Biology</i> , 2010 , 24, 1299-306	6	43
220	Measuring genetic diversity in translocation programmes: principles and application to a chimpanzee release project. <i>Animal Conservation</i> , 2002 , 5, 225-236	3.2	43
219	Domestication of cattle: Two or three events?. <i>Evolutionary Applications</i> , 2019 , 12, 123-136	4.8	42
218	Conservation Biology Framework for the Release of Wild-Born Orphaned Chimpanzees into the Konkouati Reserve, Congo. <i>Conservation Biology</i> , 2001 , 15, 1247-1257	6	42
217	A molecular phylogeny of African kestrels with reference to divergence across the Indian Ocean. <i>Molecular Phylogenetics and Evolution</i> , 2002 , 25, 267-77	4.1	41
216	Mitochondrial phylogeography and subspecific variation in the red panda (<i>Ailurus fulgens</i>): implications for conservation. <i>Molecular Phylogenetics and Evolution</i> , 2005 , 36, 78-89	4.1	41
215	Patterns of cryptic hybridization revealed using an integrative approach: a case study on genetids (Carnivora, Viverridae, <i>Genetta</i> spp.) from the southern African subregion. <i>Biological Journal of the Linnean Society</i> , 2005 , 86, 11-33	1.9	41
214	Mitochondrial phylogeography and demographic history of the vicuña: implications for conservation. <i>Heredity</i> , 2007 , 99, 70-80	3.6	40
213	Multilocus DNA fingerprints in gallinaceous birds: general approach and problems. <i>Heredity</i> , 1992 , 68 (Pt 6), 481-94	3.6	40
212	More grist for the mill? Species delimitation in the genomic era and its implications for conservation. <i>Conservation Genetics</i> , 2019 , 20, 101-113	2.6	39
211	Molecular adaptation of alanine:glyoxylate aminotransferase targeting in primates. <i>Molecular Biology and Evolution</i> , 2000 , 17, 387-400	8.3	38
210	Permanent Genetic Resources added to Molecular Ecology Resources Database 1 December 2009-31 January 2010. <i>Molecular Ecology Resources</i> , 2010 , 10, 576-9	8.4	37
209	The double origin of Iberian peninsular chameleons. <i>Biological Journal of the Linnean Society</i> , 2002 , 75, 1-7	1.9	37
208	Mating frequency and mating system of the polygynous ant, <i>Leptothorax acervorum</i> . <i>Molecular Ecology</i> , 2001 , 10, 2719-28	5.7	37

207	Invertebrate biodiversity affects predator fitness and hence potential to control pests in crops. <i>Biological Control</i> , 2009 , 51, 499-506	3.8	36
206	Accurate population size estimates are vital parameters for conserving the giant panda. <i>Ursus</i> , 2009 , 20, 56-62	1.4	35
205	DNA Fingerprinting in a Butterfly, <i>Bicyclus anynana</i> (Satyridae). <i>Journal of Heredity</i> , 1993 , 84, 195-200	2.4	35
204	Genetic structuring and recent demographic history of red pandas (<i>Ailurus fulgens</i>) inferred from microsatellite and mitochondrial DNA. <i>Molecular Ecology</i> , 2011 , 20, 2662-75	5.7	34
203	Population Genomics Reveals Low Genetic Diversity and Adaptation to Hypoxia in Snub-Nosed Monkeys. <i>Molecular Biology and Evolution</i> , 2016 , 33, 2670-81	8.3	33
202	Distinguishing gorilla mitochondrial sequences from nuclear integrations and PCR recombinants: guidelines for their diagnosis in complex sequence databases. <i>Molecular Phylogenetics and Evolution</i> , 2007 , 43, 553-66	4.1	32
201	Genets (Carnivora, Viverridae) in Africa: an evolutionary synthesis based on cytochrome b sequences and morphological characters. <i>Biological Journal of the Linnean Society</i> , 2004 , 81, 589-610	1.9	32
200	Where's the conservation in conservation genetics?. <i>Conservation Biology</i> , 2008 , 22, 802-4	6	31
199	Sistemática, taxonomía y domesticación de alpacas y llamas: nueva evidencia cromosómica y molecular. <i>Revista Chilena De Historia Natural</i> , 2007 , 80,	1.8	31
198	Quantitative evaluation of hybridization and the impact on biodiversity conservation. <i>Ecology and Evolution</i> , 2017 , 7, 320-330	2.8	30
197	Diversity, genetic structure and evidence of outcrossing in British populations of the rock fern <i>Adiantum capillus-veneris</i> using microsatellites. <i>Molecular Ecology</i> , 2001 , 10, 1881-94	5.7	30
196	Next-generation conservation genetics and biodiversity monitoring. <i>Evolutionary Applications</i> , 2018 , 11, 1029-1034	4.8	29
195	DNA identification of primate bushmeat from urban markets in Guinea-Bissau and its implications for conservation. <i>Biological Conservation</i> , 2013 , 167, 43-49	6.2	29
194	Revisiting demographic processes in cattle with genome-wide population genetic analysis. <i>Frontiers in Genetics</i> , 2015 , 6, 191	4.5	29
193	Extinctions, genetic erosion and conservation options for the black rhinoceros (<i>Diceros bicornis</i>). <i>Scientific Reports</i> , 2017 , 7, 41417	4.9	28
192	Demography and rapid local adaptation shape Creole cattle genome diversity in the tropics. <i>Evolutionary Applications</i> , 2019 , 12, 105-122	4.8	28
191	Extra-pair fertilization and effective population size in the song sparrow <i>Melospiza melodia</i> . <i>Journal of Avian Biology</i> , 2006 , 37, 572-578	1.9	28
190	Microsatellite markers for the earthworm <i>Lumbricus rubellus</i> . <i>Molecular Ecology Notes</i> , 2006 , 6, 325-327		28

189	Habitat fragmentation and genetic diversity in natural populations of the Bornean elephant: Implications for conservation. <i>Biological Conservation</i> , 2016 , 196, 80-92	6.2	27
188	Multiple introductions and environmental factors affecting the establishment of invasive species on a volcanic island. <i>Soil Biology and Biochemistry</i> , 2015 , 85, 89-100	7.5	26
187	Gastrointestinal symbionts of chimpanzees in Cantanhez National Park, Guinea-Bissau with respect to habitat fragmentation. <i>American Journal of Primatology</i> , 2013 , 75, 1032-41	2.5	26
186	Complex phylogeographic history of central African forest elephants and its implications for taxonomy. <i>BMC Evolutionary Biology</i> , 2007 , 7, 244	3	26
185	Evaluation of temperature gradient gel electrophoresis for the analysis of prey DNA within the guts of invertebrate predators. <i>Bulletin of Entomological Research</i> , 2006 , 96, 295-304	1.7	26
184	Dynamics and genetics of a disease-driven species decline to near extinction: lessons for conservation. <i>Scientific Reports</i> , 2016 , 6, 30772	4.9	26
183	The Challenges of Linking Ecosystem Services to Biodiversity. <i>Advances in Ecological Research</i> , 2016 , 54, 87-134	4.6	26
182	Fragmentation genetics of rainforest animals: insights from recent studies. <i>Conservation Genetics</i> , 2014 , 15, 245-260	2.6	25
181	Polygamy slows down population divergence in shorebirds. <i>Evolution; International Journal of Organic Evolution</i> , 2017 , 71, 1313-1326	3.8	24
180	Population transcriptomes reveal synergistic responses of DNA polymorphism and RNA expression to extreme environments on the Qinghai-Tibetan Plateau in a predatory bird. <i>Molecular Ecology</i> , 2017 , 26, 2993-3010	5.7	24
179	Exonic versus intronic SNPs: contrasting roles in revealing the population genetic differentiation of a widespread bird species. <i>Heredity</i> , 2015 , 114, 1-9	3.6	24
178	Conservation Genetic Resources for Effective Species Survival (ConGRESS): Bridging the divide between conservation research and practice. <i>Journal for Nature Conservation</i> , 2013 , 21, 433-437	2.3	24
177	Isolation and characterisation of main olfactory and vomeronasal receptor gene families from the Atlantic salmon (<i>Salmo salar</i>). <i>Gene</i> , 2006 , 371, 257-67	3.8	24
176	Population genetic structure of and inbreeding in an insular cattle breed, the Jersey, and its implications for genetic resource management. <i>Heredity</i> , 2004 , 92, 396-401	3.6	24
175	Orangutans venture out of the rainforest and into the Anthropocene. <i>Science Advances</i> , 2018 , 4, e1701422	4.3	24
174	Marine Turtles in the Turks and Caicos Islands: Remnant Rookeries, Regionally Significant Foraging Stocks, and a Major Turtle Fishery. <i>Chelonian Conservation and Biology</i> , 2009 , 8, 192-207	0.9	23
173	Mitochondrial DNA Variation and Systematics of the Guanaco (<i>Lama guanicoe</i> , Artiodactyla: Camelidae). <i>Journal of Mammalogy</i> , 2008 , 89, 269-281	1.8	23
172	Genetic composition of the Ascension Island green turtle rookery based on mitochondrial DNA: implications for sampling and diversity. <i>Endangered Species Research</i> , 2007 , 3, 145-158	2.5	23

171	Genomic analysis of the domestication and post-Spanish conquest evolution of the llama and alpaca. <i>Genome Biology</i> , 2020 , 21, 159	18.3	22
170	Nuclear DNA recapitulates the cryptic mitochondrial lineages of <i>Lumbricus rubellus</i> and suggests the existence of cryptic species in an ecotoxicological soil sentinel. <i>Biological Journal of the Linnean Society</i> , 2013 , 110, 780-795	1.9	22
169	Conservation of deer: contributions from molecular biology, evolutionary ecology, and reproductive physiology. <i>Journal of Zoology</i> , 1997 , 243, 461-484	2	22
168	Mitochondrial phylogeography and population history of finless porpoises in Sino-Japanese waters. <i>Biological Journal of the Linnean Society</i> , 2008 , 95, 193-204	1.9	22
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