Michael W. Bruford

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

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

#	Paper	IF	Citations
332	The Coalition for Conservation Genetics: Working across organizations to build capacity and achieve change in policy and practice. <i>Conservation Science and Practice</i> , 2022 , 4,	2.2	O
331	Genomic erosion in a demographically recovered bird species during conservation rescue <i>Conservation Biology</i> , 2022 , e13918	6	0
330	The influence of chalk grasslands on butterfly phenology and ecology. <i>Ecology and Evolution</i> , 2021 , 11, 14521-14539	2.8	
329	Wildlife conservation and management in China: achievements, challenges and perspectives. <i>National Science Review</i> , 2021 , 8, nwab042	10.8	5
328	Climate-driven flyway changes and memory-based long-distance migration. <i>Nature</i> , 2021 , 591, 259-264	50.4	9
327	Authors Reply to Letter to the Editor: Continued improvement to genetic diversity indicator for CBD. Conservation Genetics, 2021, 22, 533-536	2.6	3
326	Draft genome of a biparental beetle species, Lethrus apterus. <i>BMC Genomics</i> , 2021 , 22, 301	4.5	
325	Global Commitments to Conserving and Monitoring Genetic Diversity Are Now Necessary and Feasible. <i>BioScience</i> , 2021 , 71, 964-976	5.7	17
324	Effective population size remains a suitable, pragmatic indicator of genetic diversity for all species, including forest trees. <i>Biological Conservation</i> , 2021 , 253, 108906	6.2	8
323	Historical Introgression from Wild Relatives Enhanced Climatic Adaptation and Resistance to Pneumonia in Sheep. <i>Molecular Biology and Evolution</i> , 2021 , 38, 838-855	8.3	13
322	Hunting pressure is a key contributor to the impending extinction of Bornean wild cattle. <i>Endangered Species Research</i> , 2021 , 45, 225-235	2.5	О
321	Ancient and modern genomes unravel the evolutionary history of the rhinoceros family. <i>Cell</i> , 2021 , 184, 4874-4885.e16	56.2	6
320	Whole-genome resequencing of worldwide wild and domestic sheep elucidates genetic diversity, introgression and agronomically important loci. <i>Molecular Biology and Evolution</i> , 2021 ,	8.3	4
319	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
318	Recent mitochondrial lineage extinction in the critically endangered Javan rhinoceros. <i>Zoological Journal of the Linnean Society</i> , 2020 , 190, 372-383	2.4	4
317	Post-2020 goals overlook genetic diversity. <i>Science</i> , 2020 , 367, 1083-1085	33.3	64
316	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

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315	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
314	Interspecific Gene Flow and the Evolution of Specialization in Black and White Rhinoceros. <i>Molecular Biology and Evolution</i> , 2020 , 37, 3105-3117	8.3	6
313	Population differentiation and historical demography of the threatened snowy plover Charadrius nivosus (Cassin, 1858). <i>Conservation Genetics</i> , 2020 , 21, 387-404	2.6	2
312	Dispersal and genetic structure in a tropical small mammal, the Bornean tree shrew (Tupaia longipes), in a fragmented landscape along the Kinabatangan River, Sabah, Malaysia. <i>BMC Genetics</i> , 2020 , 21, 43	2.6	3
311	Estimating the population size of the Sanje mangabey (Cercocebus sanjei) using acoustic distance sampling. <i>American Journal of Primatology</i> , 2020 , 82, e23083	2.5	4
310	Paternal Origins and Migratory Episodes of Domestic Sheep. <i>Current Biology</i> , 2020 , 30, 4085-4095.e6	6.3	12
309	Set ambitious goals for biodiversity and sustainability. <i>Science</i> , 2020 , 370, 411-413	33.3	92
308	Chasing a ghost: notes on the present distribution and conservation of the sooty mangabey (Cercocebus atys) in Guinea-Bissau, West Africa. <i>Primates</i> , 2020 , 61, 357-363	1.7	
307	An ancient hybridization event reconciles mito-nuclear discordance among spiral-horned antelopes. Journal of Mammalogy, 2019 , 100, 1144-1155	1.8	2
306	Rapid identification and interpretation of gene-environment associations using the new R.SamBada landscape genomics pipeline. <i>Molecular Ecology Resources</i> , 2019 , 19, 1355-1365	8.4	8
305	Genomic selection strategies for breeding adaptation and production in dairy cattle under climate change. <i>Heredity</i> , 2019 , 123, 307-317	3.6	11
304	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
303	Domestication of cattle: Two or three events?. Evolutionary Applications, 2019, 12, 123-136	4.8	42
302	Demography and rapid local adaptation shape Creole cattle genome diversity in the tropics. <i>Evolutionary Applications</i> , 2019 , 12, 105-122	4.8	28
301	Messing about on the river: the role of geographic barriers in shaping the genetic structure of Bornean small mammals in a fragmented landscape. <i>Conservation Genetics</i> , 2019 , 20, 691-704	2.6	9
300	Inferring fine-scale spatial structure of the brown bear (Ursus arctos) population in the Carpathians prior to infrastructure development. <i>Scientific Reports</i> , 2019 , 9, 9494	4.9	8
299	21. Mitochondrial DNA Diversity In Modern Sheep: Implications For Domestication 2019 , 306-316		3
298	Rapid ecological specialization despite constant population sizes. <i>PeerJ</i> , 2019 , 7, e6476	3.1	1

297	Ecology, conservation, and phylogenetic position of the Madagascar Jacana Actophilornis albinucha. <i>Ostrich</i> , 2019 , 90, 315-326	0.9	О
296	The genomics of domestication special issue editorial. <i>Evolutionary Applications</i> , 2019 , 12, 3-5	4.8	1
295	Landscape Genetics Applied to the Conservation of Primates in Flooded Forests: A Case Study of Orangutans in the Lower Kinabatangan Wildlife Sanctuary 2019 , 297-303		
294	Genetic diversity and cryptic population re-establishment: management implications for the Bojer skink (Gongylomorphus bojerii). <i>Conservation Genetics</i> , 2019 , 20, 137-152	2.6	1
293	The hidden costs of living in a transformed habitat: Ecological and evolutionary consequences in a tripartite mutualistic system with a keystone mistletoe. <i>Science of the Total Environment</i> , 2019 , 651, 274	4 0 -2 7 4	8 ⁵
292	Genome-wide differential DNA methylation in tropically adapted Creole cattle and their Iberian ancestors. <i>Animal Genetics</i> , 2019 , 50, 15-26	2.5	16
291	Intentional Genetic Manipulation a conservation threat. Conservation Genetics Resources, 2019, 11, 237-247	0.8	12
290	Convergent genomic signatures of domestication in sheep and goats. <i>Nature Communications</i> , 2018 , 9, 813	17.4	112
289	Genetic and genomic monitoring with minimally invasive sampling methods. <i>Evolutionary Applications</i> , 2018 , 11, 1094-1119	4.8	72
288	Genetic analyses favour an ancient and natural origin of elephants on Borneo. <i>Scientific Reports</i> , 2018 , 8, 880	4.9	5
287	Transcription-Associated Mutation Promotes RNA Complexity in Highly Expressed Genes-A Major New Source of Selectable Variation. <i>Molecular Biology and Evolution</i> , 2018 , 35, 1104-1119	8.3	2
286	Next-generation metrics for monitoring genetic erosion within populations of conservation concern. <i>Evolutionary Applications</i> , 2018 , 11, 1066-1083	4.8	59
285	Comparing genetic diversity and demographic history in co-distributed wild South American camelids. <i>Heredity</i> , 2018 , 121, 387-400	3.6	19
284	Can Riparian Forest Buffers Increase Yields From Oil Palm Plantations?. <i>Earthm Future</i> , 2018 , 6, 1082-10	96 .9	O
283	Disrupted dispersal and its genetic consequences: Comparing protected and threatened baboon populations (Papio papio) in West Africa. <i>PLoS ONE</i> , 2018 , 13, e0194189	3.7	3
282	Walking in a heterogeneous landscape: Dispersal, gene flow and conservation implications for the giant panda in the Qinling Mountains. <i>Evolutionary Applications</i> , 2018 , 11, 1859-1872	4.8	12
281	Next-generation conservation genetics and biodiversity monitoring. <i>Evolutionary Applications</i> , 2018 , 11, 1029-1034	4.8	29
2 80	Primate conservation genetics 2018 , 1-2		

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279	Contrasting evolutionary history, anthropogenic declines and genetic contact in the northern and southern white rhinoceros (). <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2018 , 285,	4.4	11
278	Altitudinal ranging of the Guizhou golden monkey (Rhinopithecus brelichi): Patterns of habitat selection and habitat use. <i>Global Ecology and Conservation</i> , 2018 , 16, e00473	2.8	
277	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,	7.6	14
276	Contrasting Patterns of Genomic Diversity Reveal Accelerated Genetic Drift but Reduced Directional Selection on X-Chromosome in Wild and Domestic Sheep Species. <i>Genome Biology and Evolution</i> , 2018 , 10, 1282-1297	3.9	15
275	Orangutans venture out of the rainforest and into the Anthropocene. Science Advances, 2018, 4, e17014	4 22 .3	24
274	The Value of Ecosystem Services from Giant Panda Reserves. <i>Current Biology</i> , 2018 , 28, 2174-2180.e7	6.3	75
273	Quantitative evaluation of hybridization and the impact on biodiversity conservation. <i>Ecology and Evolution</i> , 2017 , 7, 320-330	2.8	30
272	Polygamy slows down population divergence in shorebirds. <i>Evolution; International Journal of Organic Evolution</i> , 2017 , 71, 1313-1326	3.8	24
271	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
270	Extinctions, genetic erosion and conservation options for the black rhinoceros (Diceros bicornis). <i>Scientific Reports</i> , 2017 , 7, 41417	4.9	28
269	Modification of river meandering by tropical deforestation. <i>Geology</i> , 2017 , 45, 511-514	5	46
268	Monitoring Changes in Genetic Diversity 2017 , 107-128		15
267	Odour dialects among wild mammals. <i>Scientific Reports</i> , 2017 , 7, 13593	4.9	3
266	Genomic signatures of adaptive introgression from European mouflon into domestic sheep. <i>Scientific Reports</i> , 2017 , 7, 7623	4.9	49
265	Genetics and Primate Conservation 2017 , 1-6		1
264	Enhancing capacity for freshwater conservation at the genetic level: a demonstration using three stream macroinvertebrates. <i>Aquatic Conservation: Marine and Freshwater Ecosystems</i> , 2017 , 27, 452-461	2.6	7
263	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
262	Dispersal of green turtles from Africal largest rookery assessed through genetic markers. <i>Marine Ecology - Progress Series</i> , 2017 , 569, 215-225	2.6	12

261	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
260	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
259	Evidence for deleterious effects of harness-mounted satellite transmitters on Saker Falcons Falco cherrug. <i>Bird Study</i> , 2016 , 63, 96-106	0.7	10
258	Genetic consequences of human forest exploitation in two colobus monkeys in Guinea Bissau. <i>Biological Conservation</i> , 2016 , 194, 194-208	6.2	4
257	Assessing Genetic Structure in Common but Ecologically Distinct Carnivores: The Stone Marten and Red Fox. <i>PLoS ONE</i> , 2016 , 11, e0145165	3.7	11
256	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
255	Landscape determinants of fine-scale genetic structure of a small rodent in a heterogeneous landscape (Hluhluwe-iMfolozi Park, South Africa). <i>Scientific Reports</i> , 2016 , 6, 29168	4.9	17
254	Non-invasive genetic identification confirms the presence of the Endangered okapi Okapia johnstoni south-west of the Congo River. <i>Oryx</i> , 2016 , 50, 134-137	1.5	5
253	Colonization of the Scottish islands via long-distance Neolithic transport of red deer (Cervus elaphus). <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2016 , 283,	4.4	15
252	The Challenges of Linking Ecosystem Services to Biodiversity. <i>Advances in Ecological Research</i> , 2016 , 54, 87-134	4.6	26
251	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
250	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
249	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
248	Kinship and Intragroup Social Dynamics in Two Sympatric African Colobus Species. <i>International Journal of Primatology</i> , 2015 , 36, 871-886	2	2
247	The role of density and relatedness in wild juvenile Atlantic salmon growth. <i>Journal of Zoology</i> , 2015 , 295, 56-64	2	6
246	Genomics and the challenging translation into conservation practice. <i>Trends in Ecology and Evolution</i> , 2015 , 30, 78-87	10.9	335
245	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
244	Contrasting genetic diversity and population structure among three sympatric Madagascan shorebirds: parallels with rarity, endemism, and dispersal. <i>Ecology and Evolution</i> , 2015 , 5, 997-1010	2.8	20

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243	Revisiting demographic processes in cattle with genome-wide population genetic analysis. <i>Frontiers in Genetics</i> , 2015 , 6, 191	4.5	29
242	Prospects and challenges for the conservation of farm animal genomic resources, 2015-2025. <i>Frontiers in Genetics</i> , 2015 , 6, 314	4.5	50
241	Genetic structure of captive and free-ranging okapi (Okapia johnstoni) with implications for management. <i>Conservation Genetics</i> , 2015 , 16, 1115-1126	2.6	5
240	Evolution and Conservation of Central African Biodiversity: Priorities for Future Research and Education in the Congo Basin and Gulf of Guinea. <i>Biotropica</i> , 2015 , 47, 6-17	2.3	9
239	Enhancing knowledge of an endangered and elusive species, the okapi, using non-invasive genetic techniques. <i>Journal of Zoology</i> , 2015 , 295, 233-242	2	3
238	Fragmentation genetics of rainforest animals: insights from recent studies. <i>Conservation Genetics</i> , 2014 , 15, 245-260	2.6	25
237	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
236	Identifying biochemical phenotypic differences between cryptic species. <i>Biology Letters</i> , 2014 , 10,	3.6	12
235	Mitochondrial DNA and morphological variation in the sentinel earthworm species Lumbricus rubellus. <i>European Journal of Soil Biology</i> , 2014 , 64, 23-29	2.9	15
234	Contrasting genetic structure of the Eurasian otter (Lutra lutra) across a latitudinal divide. <i>Journal of Mammalogy</i> , 2014 , 95, 814-823	1.8	5
233	A horizon scan for species conservation by zoos and aquariums. Zoo Biology, 2014, 33, 375-80	1.6	12
232	Assessing the impact of hunting pressure on population structure of Guinea baboons (Papio papio) in Guinea-Bissau. <i>Conservation Genetics</i> , 2014 , 15, 1339-1355	2.6	14
231	Assessing the spatial dependence of adaptive loci in 43 European and Western Asian goat breeds using AFLP markers. <i>PLoS ONE</i> , 2014 , 9, e86668	3.7	12
230	Distinct and diverse: range-wide phylogeography reveals ancient lineages and high genetic variation in the endangered okapi (Okapia johnstoni). <i>PLoS ONE</i> , 2014 , 9, e101081	3.7	13
229	Admixture analysis in relation to pedigree studies of introgression in a minority British cattle breed: the Lincoln Red. <i>Journal of Animal Breeding and Genetics</i> , 2014 , 131, 19-26	2.9	1
228	Whole-genome analyses resolve early branches in the tree of life of modern birds. <i>Science</i> , 2014 , 346, 1320-31	33.3	1182
227	Comparative genomics reveals insights into avian genome evolution and adaptation. <i>Science</i> , 2014 , 346, 1311-20	33.3	628
226	Comparative evaluation of potential indicators and temporal sampling protocols for monitoring genetic erosion. <i>Evolutionary Applications</i> , 2014 , 7, 984-98	4.8	77

225	Genetic evidence for spatio-temporal changes in the dispersal patterns of two sympatric African colobine monkeys. <i>American Journal of Physical Anthropology</i> , 2013 , 150, 464-74	2.5	10
224	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
223	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
222	The genetic legacy of the 19th-century decline of the British polecat: evidence for extensive introgression from feral ferrets. <i>Molecular Ecology</i> , 2013 , 22, 5130-47	5.7	16
221	Ecology. Essential biodiversity variables. <i>Science</i> , 2013 , 339, 277-8	33.3	809
220	Demographic loss, genetic structure and the conservation implications for Indian tigers. Proceedings of the Royal Society B: Biological Sciences, 2013, 280, 20130496	4.4	59
219	Intraguild predation in winter wheat: prey choice by a common epigeal carabid consuming spiders. Journal of Applied Ecology, 2013 , 50, 271-279	5.8	55
218	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
217	Bringing genetic diversity to the forefront of conservation policy and management. <i>Conservation Genetics Resources</i> , 2013 , 5, 593-598	0.8	91
216	DNA sequence variation and methylation in an arsenic tolerant earthworm population. <i>Soil Biology and Biochemistry</i> , 2013 , 57, 524-532	7.5	58
215	Genetic consequences of historical anthropogenic and ecological events on giant pandas. <i>Ecology</i> , 2013 , 94, 2346-57	4.6	54
214	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
213	Nuclear DNA recapitulates the cryptic mitochondrial lineages of Lumbricus rubellusand suggests the existence of cryptic species in an ecotoxological soil sentinel. <i>Biological Journal of the Linnean Society</i> , 2013 , 110, 780-795	1.9	22
212	Missense SNP of the MC1R gene is associated with plumage variation in the Gyrfalcon (Falco rusticolus). <i>Animal Genetics</i> , 2012 , 43, 460-2	2.5	14
211	Molecular tools and analytical approaches for the characterization of farm animal genetic diversity. <i>Animal Genetics</i> , 2012 , 43, 483-502	2.5	82
210	Rapid ongoing decline of Baird's tapir in Cusuco National Park, Honduras. <i>Integrative Zoology</i> , 2012 , 7, 420-428	1.9	6
209	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
208	Effective population size dynamics and the demographic collapse of Bornean orang-utans. <i>PLoS ONE</i> , 2012 , 7, e49429	3.7	62

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207	A panel of microsatellite markers for genetic studies of European polecats (Mustela putorius) and ferrets (Mustela furo). <i>European Journal of Wildlife Research</i> , 2012 , 58, 629-633	2	3
206	Biodiversity and conservation genetics research in Central Africa: new approaches and avenues for international collaboration. <i>Conservation Genetics Resources</i> , 2012 , 4, 523-525	0.8	5
205	First record of Neoxysomatium brevicaudatum through the non-invasive sampling of Anguis fragilis: complementary morphological and molecular detection. <i>Journal of Helminthology</i> , 2012 , 86, 125-9	1.6	11
204	Morphometric differentiation of Tetramesa leucospae Zerova & Madjdzadeh, 2005, populations associated with two geographically isolated grass species in Iran. <i>Zoology in the Middle East</i> , 2011 , 52, 79-88	0.7	2
203	Genetic diversity of sheep breeds from Albania, Greece, and Italy assessed by mitochondrial DNA and nuclear polymorphisms (SNPs). <i>Scientific World Journal, The</i> , 2011 , 11, 1641-59	2.2	17
202	Primate conservation: measuring and mitigating trade in primates. <i>Endangered Species Research</i> , 2011 , 13, 159-161	2.5	75
201	Bayesian clustering techniques and progressive partitioning to identify population structuring within a recovering otter population in the UK. <i>Journal of Applied Ecology</i> , 2011 , 48, 1206-1217	5.8	18
200	Genetic structuring and recent demographic history of red pandas (Ailurus fulgens) inferred from microsatellite and mitochondrial DNA. <i>Molecular Ecology</i> , 2011 , 20, 2662-75	5.7	34
199	Molecular evidence for Pleistocene refugia at the eastern edge of the Tibetan Plateau. <i>Molecular Ecology</i> , 2011 , 20, 3014-26	5.7	53
198	Lack of Evidence of Simian Immunodeficiency Virus Infection Among Nonhuman Primates in Ta [®] National Park, Cle d'Ivoire: Limitations of Noninvasive Methods and SIV Diagnostic Tools for Studies of Primate Retroviruses. <i>International Journal of Primatology</i> , 2011 , 32, 288-307	2	5
197	The use of non-invasive molecular techniques to confirm the presence of mountain bongo Tragelaphus eurycerus isaaci populations in Kenya and preliminary inference of their mitochondrial genetic variation. <i>Conservation Genetics</i> , 2011 , 12, 745-751	2.6	11
196	Substantial molecular variation and low genetic structure in Kenyall black rhinoceros: implications for conservation. <i>Conservation Genetics</i> , 2011 , 12, 1575-1588	2.6	10
195	Genotyping faeces of red pandas (Ailurus fulgens): implications for population estimation. <i>European Journal of Wildlife Research</i> , 2011 , 57, 1231-1235	2	3
194	Promoting collaboration between livestock and wildlife conservation genetics communities. <i>Conservation Genetics Resources</i> , 2011 , 3, 785-788	0.8	6
193	Mitochondrial DNA monomorphism in Red-billed Choughs Pyrrhocorax pyrrhocorax in the United Kingdom. <i>Bird Study</i> , 2011 , 58, 213-216	0.7	1
192	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
191	Population structure in the South American tern Sterna hirundinacea in the South Atlantic: two populations with distinct breeding phenologies. <i>Journal of Avian Biology</i> , 2010 , 41, 378-387	1.9	19
190	Unravelling migratory connectivity in marine turtles using multiple methods. <i>Journal of Applied Ecology</i> , 2010 , 47, 769-778	5.8	71

189	The sequence and de novo assembly of the giant panda genome. <i>Nature</i> , 2010 , 463, 311-7	50.4	864
188	Spatial Trends of Genetic Variation of Domestic Ruminants in Europe. <i>Diversity</i> , 2010 , 2, 932-945	2.5	19
187	2BAD: an application to estimate the parental contributions during two independent admixture events. <i>Molecular Ecology Resources</i> , 2010 , 10, 538-41	8.4	10
186	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
185	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
184	Cryptic sexual size dimorphism in Malagasy plovers Charadrius spp Ostrich, 2010 , 81, 173-178	0.9	6
183	Microsatellite variation and significant population genetic structure of endangered finless porpoises (Neophocaena phocaenoides) in Chinese coastal waters and the Yangtze River. <i>Marine Biology</i> , 2010 , 157, 1453-1462	2.5	15
182	PCR primers for microsatellite loci in a Madagascan waterbird, the Sakalava Rail (Amaurornis olivieri). <i>Conservation Genetics Resources</i> , 2010 , 2, 273-277	0.8	
181	Isolation and characterization of 13 tetranucleotide microsatellite loci in the Stone marten (Martes foina). <i>Conservation Genetics Resources</i> , 2010 , 2, 317-319	0.8	13
180	Microsatellite loci for the okapi (Okapia johnstoni). <i>Conservation Genetics Resources</i> , 2010 , 2, 337-339	0.8	4
179	Microsatellite markers for the proboscis monkey (Nasalis larvatus). <i>Conservation Genetics Resources</i> , 2010 , 2, 159-163	0.8	7
178	Genetic structure of the Black Bog Ant (Formica picea Nylander) in the United Kingdom. <i>Conservation Genetics</i> , 2010 , 11, 823-834	2.6	4
177	A new method for quantifying genotyping errors for noninvasive genetic studies. <i>Conservation Genetics</i> , 2010 , 11, 1567-1571	2.6	16
176	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
175	A large panel of microsatellite markers for genetic studies in the infra-order catarrhini. <i>Folia Primatologica</i> , 2009 , 80, 63-9	1.2	11
174	Minisatellite DNA markers in the chicken genome. <i>Animal Genetics</i> , 2009 , 25, 381-389	2.5	5
173	Invertebrate biodiversity affects predator fitness and hence potential to control pests in crops. <i>Biological Control</i> , 2009 , 51, 499-506	3.8	36
172	Mitochondrial genetic diversity and structure of the European otter (Lutra lutra) in Britain. <i>Conservation Genetics</i> , 2009 , 10, 733-737	2.6	17

(2008-2009)

171	Isolation and characterisation of 11 tetranucleotide microsatellite loci in the common genet (Genetta genetta). <i>Conservation Genetics</i> , 2009 , 10, 1931-1934	2.6	1
170	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
169	Turtle groups or turtle soup: dispersal patterns of hawksbill turtles in the Caribbean. <i>Molecular Ecology</i> , 2009 , 18, 4841-53	5.7	88
168	Molecular structure in peripheral dog breeds: Portuguese native breeds as a case study. <i>Animal Genetics</i> , 2009 , 40, 383-92	2.5	9
167	The population genetic effects of ancestry and admixture in a subdivided cattle breed. <i>Animal Genetics</i> , 2009 , 40, 393-400	2.5	19
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