

Nuno Ferrand

List of Publications by Year in descending order

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142
papers

6,426
citations

66234

42
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79541

73
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146
docs citations

146
times ranked

7166
citing authors

#	ARTICLE	IF	CITATIONS
1	Genetic differentiation in Eurasian Woodcock (<i>Scolopax rusticola</i>) from the Azores. <i>Ibis</i> , 2022, 164, 313-319.	1.0	1
2	African climate and geomorphology drive evolution and ghost introgression in sable antelope. <i>Molecular Ecology</i> , 2022, 31, 2968-2984.	2.0	8
3	A loss-of-function mutation in RORB disrupts saltatorial locomotion in rabbits. <i>PLoS Genetics</i> , 2021, 17, e1009429.	1.5	10
4	Evolutionary history of the roan antelope across its African range. <i>Journal of Biogeography</i> , 2021, 48, 2812-2827.	1.4	4
5	De novo whole-genome assembly and resequencing resources for the roan (<i>Hippotragus</i>)	0.8	4
6	Pervasive hybridization with local wild relatives in Western European grapevine varieties. <i>Science Advances</i> , 2021, 7, eabi8584.	4.7	11
7	The Global Museum: natural history collections and the future of evolutionary science and public education. <i>PeerJ</i> , 2020, 8, e8225.	0.9	81
8	Genetic characterization of green turtles (<i>Chelonia mydas</i>) from São Tomé and Príncipe: Insights on species recruitment and dispersal in the Gulf of Guinea. <i>Journal of Experimental Marine Biology and Ecology</i> , 2019, 518, 151181.	0.7	6
9	Whole Genome Sequencing and Re-sequencing of the Sable Antelope (<i>Hippotragus niger</i>): A Resource for Monitoring Diversity in ex Situ and in Situ Populations. <i>G3: Genes, Genomes, Genetics</i> , 2019, 9, 1785-1793.	0.8	18
10	Unravelling population processes over the Late Pleistocene driving contemporary genetic divergence in Palearctic buzzards. <i>Molecular Phylogenetics and Evolution</i> , 2019, 134, 269-281.	1.2	8
11	Parallel adaptation of rabbit populations to myxoma virus. <i>Science</i> , 2019, 363, 1319-1326.	6.0	124
12	Angolan Biodiversity: Towards a Modern Synthesis. , 2019, , 3-14.		5
13	Conclusions: Biodiversity Research and Conservation Opportunities. , 2019, , 543-549.		3
14	Mitochondrial phylogeography of the Iberian endemic frog <i>Rana iberica</i> , with implications for its conservation. <i>Environmental Epigenetics</i> , 2018, 64, 755-764.	0.9	12
15	A genomic map of clinal variation across the European rabbit hybrid zone. <i>Molecular Ecology</i> , 2018, 27, 1457-1478.	2.0	30
16	The curious case of <i>Bradypus variegatus</i> sloths: populations in threatened habitats are biodiversity components needing protection. <i>Biodiversity and Conservation</i> , 2018, 27, 1291-1308.	1.2	4
17	Cryptic population structure reveals low dispersal in Iberian wolves. <i>Scientific Reports</i> , 2018, 8, 14108.	1.6	36
18	Changes in brain architecture are consistent with altered fear processing in domestic rabbits. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 7380-7385.	3.3	45

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19	Draft genome assembly of the invasive cane toad, <i>Rhinella marina</i> . <i>GigaScience</i> , 2018, 7, .	3.3	60
20	The roles of allopatric fragmentation and niche divergence in intraspecific lineage diversification in the common midwife toad (<i>Alytes obstetricans</i>). <i>Journal of Biogeography</i> , 2018, 45, 2146-2158.	1.4	24
21	Dwarfism and Altered Craniofacial Development in Rabbits Is Caused by a 12.1 kb Deletion at the <i>HMGA2</i> Locus. <i>Genetics</i> , 2017, 205, 955-965.	1.2	30
22	Hybridization following population collapse in a critically endangered antelope. <i>Scientific Reports</i> , 2016, 6, 18788.	1.6	35
23	Candidate genes underlying heritable differences in reproductive seasonality between wild and domestic rabbits. <i>Animal Genetics</i> , 2015, 46, 418-425.	0.6	14
24	Understanding the mechanisms of antitropical divergence in the seabird <i>Stercorarius pomarinus</i> using a multilocus approach. <i>Molecular Ecology</i> , 2015, 24, 3122-3137.	2.0	15
25	Levels and Patterns of Genetic Diversity and Population Structure in Domestic Rabbits. <i>PLoS ONE</i> , 2015, 10, e0144687.	1.1	38
26	Genealogy of the nuclear β -fibrinogen intron 7 in <i>Lissotriton boscai</i> (Caudata, Salamandridae): concordance with mtDNA and implications for phylogeography and speciation. <i>Contributions To Zoology</i> , 2015, 84, 193-215.	0.2	18
27	Low persistence in nature of captive reared rabbits after restocking operations. <i>European Journal of Wildlife Research</i> , 2015, 61, 591-599.	0.7	9
28	First estimates of genetic diversity for the highly endangered giant sable antelope using a set of 57 microsatellites. <i>European Journal of Wildlife Research</i> , 2015, 61, 313-317.	0.7	10
29	Multilocus phylogeography of the common midwife toad, <i>Alytes obstetricans</i> (Anura, Alytidae): Contrasting patterns of lineage diversification and genetic structure in the Iberian refugium. <i>Molecular Phylogenetics and Evolution</i> , 2015, 93, 363-379.	1.2	27
30	Limited gene flow and high genetic diversity in the threatened Betic midwife toad (<i>Alytes dickhilleni</i>): evolutionary and conservation implications. <i>Conservation Genetics</i> , 2015, 16, 459-476.	0.8	11
31	Molecular evidence for cryptic candidate species in Iberian <i>Pelodytes</i> (Anura, Pelodytidae). <i>Molecular Phylogenetics and Evolution</i> , 2015, 83, 224-241.	1.2	22
32	Real-time assessment of hybridization between wolves and dogs: combining noninvasive samples with ancestry informative markers. <i>Molecular Ecology Resources</i> , 2015, 15, 317-328.	2.2	53
33	The Genomic Architecture of Population Divergence between Subspecies of the European Rabbit. <i>PLoS Genetics</i> , 2014, 10, e1003519.	1.5	82
34	Intraspecific genetic variation in the common midwife toad (<i>Alytes obstetricans</i>): subspecies assignment using mitochondrial and microsatellite markers. <i>Journal of Zoological Systematics and Evolutionary Research</i> , 2014, 52, 170-175.	0.6	13
35	Rabbit genome analysis reveals a polygenic basis for phenotypic change during domestication. <i>Science</i> , 2014, 345, 1074-1079.	6.0	343
36	Multilocus assessment of phylogenetic relationships in <i>Alytes</i> (Anura, Alytidae). <i>Molecular Phylogenetics and Evolution</i> , 2014, 79, 270-278.	1.2	23

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37	Colonization history of Mallorca Island by the European rabbit, <i>Oryctolagus cuniculus</i> , and the Iberian hare, <i>Lepus granatensis</i> (Lagomorpha: Leporidae). <i>Biological Journal of the Linnean Society</i> , 2014, 111, 748-760.	0.7	7
38	Steep clines within a highly permeable genome across a hybrid zone between two subspecies of the European rabbit. <i>Molecular Ecology</i> , 2013, 22, 2511-2525.	2.0	44
39	Identifying Loci Under Selection Against Gene Flow in Isolation-with-Migration Models. <i>Genetics</i> , 2013, 194, 211-233.	1.2	58
40	A Comparison of Brain Gene Expression Levels in Domesticated and Wild Animals. <i>PLoS Genetics</i> , 2012, 8, e1002962.	1.5	130
41	Evidence for Widespread Positive and Purifying Selection Across the European Rabbit (<i>Oryctolagus</i>)	3.5	71
42	Copy number polymorphism in the β -globin gene cluster of European rabbit (<i>Oryctolagus cuniculus</i>). <i>Heredity</i> , 2012, 108, 531-536.	1.2	4
43	Genetic variability in mitochondrial and nuclear genes of <i>Larus dominicanus</i> (Charadriiformes)	0.6	8
44	Conservation genetics of the endangered Dorcas gazelle (<i>Gazella dorcas</i> spp.) in Northwestern Africa. <i>Conservation Genetics</i> , 2012, 13, 1003-1015.	0.8	26
45	Integrating molecular ecology and predictive modelling: implications for the conservation of the barbastelle bat (<i>Barbastella barbastellus</i>) in Portugal. <i>European Journal of Wildlife Research</i> , 2012, 58, 721-732.	0.7	5
46	Divide to conquer: a complex pattern of biodiversity depicted by vertebrate components in the Brazilian Atlantic Forest. <i>Biological Journal of the Linnean Society</i> , 2012, 107, 39-55.	0.7	40
47	Postglacial colonization of Europe by the barbastelle bat: agreement between molecular data and past predictive modelling. <i>Molecular Ecology</i> , 2012, 21, 2761-2774.	2.0	37
48	The Genetic Structure of Domestic Rabbits. <i>Molecular Biology and Evolution</i> , 2011, 28, 1801-1816.	3.5	101
49	Hybridization and massive mtDNA unidirectional introgression between the closely related Neotropical toads <i>Rhinella marina</i> and <i>R. schneideri</i> inferred from mtDNA and nuclear markers. <i>BMC Evolutionary Biology</i> , 2011, 11, 264.	3.2	70
50	Permanent Genetic Resources added to Molecular Ecology Resources Database 1 December 2010–31 January 2011. <i>Molecular Ecology Resources</i> , 2011, 11, 586-589.	2.2	38
51	Genetic evidence for multiple events of hybridization between wolves and domestic dogs in the Iberian Peninsula. <i>Molecular Ecology</i> , 2011, 20, 5154-5166.	2.0	118
52	INTERSPECIFIC X-CHROMOSOME AND MITOCHONDRIAL DNA INTROGRESSION IN THE IBERIAN HARE: SELECTION OR ALLELE SURFING?. <i>Evolution; International Journal of Organic Evolution</i> , 2011, 65, 1956-1968.	1.1	29
53	Recent evolutionary history of the Iberian endemic lizards <i>Podarcis bocagei</i> (Seoane, 1884) and <i>Podarcis carbonelli</i> Pérez-Mellado, 1981 (Squamata: Lacertidae) revealed by allozyme and microsatellite markers. <i>Zoological Journal of the Linnean Society</i> , 2011, 162, 184-200.	1.0	13
54	New primers for the amplification and sequencing of nuclear loci in a taxonomically wide set of reptiles and amphibians. <i>Conservation Genetics Resources</i> , 2010, 2, 181-185.	0.4	57

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55	SPECIATION IN THE EUROPEAN RABBIT (<i>ORYCTOLAGUS CUNICULUS</i>): ISLANDS OF DIFFERENTIATION ON THE X CHROMOSOME AND AUTOSOMES. <i>Evolution; International Journal of Organic Evolution</i> , 2010, 64, 3443-3460.	1.1	71
56	Extensive Gene Conversion Drives the Concerted Evolution of Paralogous Copies of the SRY Gene in European Rabbits. <i>Molecular Biology and Evolution</i> , 2010, 27, 2437-2440.	3.5	26
57	Evolutionary and functional insights into the mechanism underlying high-altitude adaptation of deer mouse hemoglobin. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 14450-14455.	3.3	202
58	The genomic legacy from the extinct <i>Lepus timidus</i> to the three hare species of Iberia: contrast between mtDNA, sex chromosomes and autosomes. <i>Molecular Ecology</i> , 2009, 18, 2643-2658.	2.0	69
59	High levels of population subdivision in a morphologically conserved Mediterranean toad (<i>Alytes</i>) nuclear genealogies. <i>Molecular Ecology</i> , 2009, 18, 5143-5160.	2.0	51
60	Genetic admixture between the Iberian endemic lizards <i>Podarcis bocagei</i> and <i>Podarcis carbonelli</i> : evidence for limited natural hybridization and a bimodal hybrid zone. <i>Journal of Zoological Systematics and Evolutionary Research</i> , 2009, 47, 368-377.	0.6	34
61	Recombination and Speciation: Loci Near Centromeres Are More Differentiated Than Loci Near Telomeres Between Subspecies of the European Rabbit (<i>Oryctolagus cuniculus</i>). <i>Genetics</i> , 2009, 181, 593-606.	1.2	92
62	Rabbit. , 2009, , 165-230.		5
63	Development of new nuclear markers and characterization of single nucleotide polymorphisms in kelp gull (<i>Larus dominicanus</i>). <i>Molecular Ecology Resources</i> , 2009, 9, 1159-1161.	2.2	6
64	Molecular analysis of hybridisation between wild and domestic cats (<i>Felis silvestris</i>) in Portugal: implications for conservation. <i>Conservation Genetics</i> , 2008, 9, 1-11.	0.8	100
65	Evidence for genetic similarity of two allopatric European hares (<i>Lepus corsicanus</i> and <i>L.</i>) 1191-1197.	1.2	39
66	Evidence for contrasting modes of selection at interacting globin genes in the European rabbit (<i>Oryctolagus cuniculus</i>). <i>Heredity</i> , 2008, 100, 602-609.	1.2	5
67	The limits of mtDNA phylogeography: complex patterns of population history in a highly structured Iberian lizard are only revealed by the use of nuclear markers. <i>Molecular Ecology</i> , 2008, 17, 4670-4683.	2.0	100
68	Reduced introgression of the Y chromosome between subspecies of the European rabbit (<i>Oryctolagus cuniculus</i>) in the Iberian Peninsula. <i>Molecular Ecology</i> , 2008, 17, 4489-4499.	2.0	45
69	Non-equilibrium estimates of gene flow inferred from nuclear genealogies suggest that Iberian and North African wall lizards (<i>Podarcis</i> spp.) are an assemblage of incipient species. <i>BMC Evolutionary Biology</i> , 2008, 8, 63.	3.2	78
70	Inferring the Evolutionary History of the European Rabbit (<i>Oryctolagus cuniculus</i>) from Molecular Markers. , 2008, , 47-63.		20
71	Overview of Lagomorph Research: What we have learned and what we still need to do. , 2008, , 381-391.		8
72	Assignment tests applied to relocate individuals of unknown origin in a threatened species, the European pond turtle (<i>Emys orbicularis</i>). <i>Amphibia - Reptilia</i> , 2007, 28, 475-484.	0.1	16

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73	Geographical variation in the golden-striped salamander, <i>Chioglossa lusitanica</i> Bocage, 1864 and the description of a newly recognized subspecies. <i>Journal of Natural History</i> , 2007, 41, 925-936.	0.2	8
74	The Molecular Basis of High-Altitude Adaptation in Deer Mice. <i>PLoS Genetics</i> , 2007, 3, e45.	1.5	173
75	Extensive intragenic recombination and patterns of linkage disequilibrium at the CSN3 locus in European rabbit. <i>Genetics Selection Evolution</i> , 2007, 39, 341-352.	1.2	7
76	The evolutionary history of the European rabbit (<i>Oryctolagus cuniculus</i>): major patterns of population differentiation and geographic expansion inferred from protein polymorphism. , 2007, , 207-235.		30
77	Patterns of hemoglobin polymorphism [β^1 -globin (HBA) and β^2 -globin (HBB)] across the contact zone of two distinct phylogeographic lineages of the European rabbit (<i>Oryctolagus cuniculus</i>). , 2007, , 237-255.		6
78	Historical biogeography and conservation of the golden-striped salamander (<i>Chioglossa lusitanica</i>) in northwestern Iberia: integrating ecological, phenotypic and phylogeographic data. , 2007, , 189-205.		11
79	Current perspectives in phylogeography and the significance of South European refugia in the creation and maintenance of European biodiversity. , 2007, , 341-357.		20
80	Conflicting phylogenetic signal of nuclear vs mitochondrial DNA markers in midwife toads (<i>Anura</i>). <i>Trends in Ecology & Evolution</i> , 2007, 44, 494-500.	1.2	45
81	Comparing patterns of nuclear and mitochondrial divergence in a cryptic species complex: the case of Iberian and North African wall lizards (<i>Podarcis</i> , Lacertidae). <i>Biological Journal of the Linnean Society</i> , 2007, 91, 121-133.	0.7	67
82	Contrasting patterns of population subdivision and historical demography in three western Mediterranean lizard species inferred from mitochondrial DNA variation. <i>Molecular Ecology</i> , 2007, 16, 1191-1205.	2.0	74
83	Isolation and characterization of microsatellite markers in pangolins (<i>Mammalia</i>). <i>Trends in Ecology & Evolution</i> , 2007, 22, 342-347.	1.7	50
84	Extensive intragenic recombination and patterns of linkage disequilibrium at the CSN3 locus in European rabbit. <i>Genetics Selection Evolution</i> , 2007, 39, 341.	1.2	5
85	Contrasting Patterns of Introgression at X-Linked Loci Across the Hybrid Zone Between Subspecies of the European Rabbit (<i>Oryctolagus cuniculus</i>). <i>Genetics</i> , 2006, 173, 919-933.	1.2	89
86	Isolation of polymorphic microsatellite loci from Eurasian woodcock (<i>Scolopax rusticola</i>) and their cross-utility in related species. <i>Molecular Ecology Notes</i> , 2006, 7, 130-132.	1.7	4
87	Extensive intraspecific polymorphism detected by SSCP at the nuclear C-mos gene in the endemic Iberian lizard <i>Lacerta schreiberi</i> . <i>Molecular Ecology</i> , 2006, 15, 731-738.	2.0	39
88	The rise and fall of the mountain hare (<i>Lepus timidus</i>) during Pleistocene glaciations: expansion and retreat with hybridization in the Iberian Peninsula. <i>Molecular Ecology</i> , 2006, 16, 605-618.	2.0	95
89	Genealogy of the nuclear β^2 -fibrinogen locus in a highly structured lizard species: comparison with mtDNA and evidence for intragenic recombination in the hybrid zone. <i>Heredity</i> , 2006, 96, 454-463.	1.2	38
90	Reexamination of the Iberian and North African <i>Podarcis</i> (<i>Squamata</i> : Lacertidae) phylogeny based on increased mitochondrial DNA sequencing. <i>Molecular Phylogenetics and Evolution</i> , 2006, 38, 266-273.	1.2	85

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91	Hares on thin ice: Introgression of mitochondrial DNA in hares and its implications for recent phylogenetic analyses. <i>Molecular Phylogenetics and Evolution</i> , 2006, 40, 640-641.	1.2	40
92	Genetic variation at chemokine receptor CCR5 in leporids: alteration at the 2nd extracellular domain by gene conversion with CCR2 in <i>Oryctolagus</i> , but not in <i>Sylvilagus</i> and <i>Lepus</i> species. <i>Immunogenetics</i> , 2006, 58, 494-501.	1.2	38
93	A 7-bp insertion in the 3' untranslated region suggests the duplication and concerted evolution of the rabbit SRY gene. <i>Genetics Selection Evolution</i> , 2006, 38, 313.	1.2	17
94	Assessing the phylogenetic signal of the nuclear β -Fibrinogen intron 7 in salamandrids (Amphibia: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 542 T	0.1	28
95	The origin of European cattle: Evidence from modern and ancient DNA. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 8113-8118.	3.3	271
96	Isolation and characterization of seven microsatellite loci in <i>Chioglossa lusitanica</i> (Urodela: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 542 T	1.7	3
97	Isolation and characterization of two dinucleotide and four tetranucleotide polymorphic microsatellite loci in the Iberian midwife toad <i>Alytes cisternasii</i> . <i>Molecular Ecology Notes</i> , 2005, 5, 767-769.	1.7	0
98	Invasion from the cold past: extensive introgression of mountain hare (<i>Lepus timidus</i>) mitochondrial DNA into three other hare species in northern Iberia. <i>Molecular Ecology</i> , 2005, 14, 2459-2464.	2.0	183
99	High levels of nucleotide diversity in the European rabbit (<i>Oryctolagus cuniculus</i>) SRY gene. <i>Animal Genetics</i> , 2005, 36, 349-351.	0.6	34
100	The evolution of the immunoglobulin heavy chain variable region (IgV H) in Leporids: an unusual case of transspecies polymorphism. <i>Immunogenetics</i> , 2005, 57, 874-882.	1.2	31
101	Phylogeny and evolution of the green lizards, <i>Lacerta</i> spp. (Squamata: Lacertidae) based on mitochondrial and nuclear DNA sequences. <i>Amphibia - Reptilia</i> , 2005, 26, 271-285.	0.1	67
102	Allelic Variation at the VHa Locus in Natural Populations of Rabbit (<i>Oryctolagus cuniculus</i> , L.). <i>Journal of Immunology</i> , 2004, 172, 1044-1053.	0.4	28
103	Genetic exchange across a hybrid zone within the Iberian endemic golden-striped salamander, <i>Chioglossa lusitanica</i> . <i>Molecular Ecology</i> , 2004, 14, 245-254.	2.0	52
104	Isolation and characterization of nine microsatellite loci in <i>Podarcis bocagei</i> (Squamata: Lacertidae). <i>Molecular Ecology Notes</i> , 2004, 4, 286-288.	1.7	21
105	Twenty polymorphic microsatellites in two of North Africa's most threatened ungulates: <i>Gazella dorcas</i> and <i>Ammotragus lervia</i> (Bovidae; Artiodactyla). <i>Molecular Ecology Notes</i> , 2004, 4, 452-455.	1.7	19
106	African Origins of the Domestic Donkey. <i>Science</i> , 2004, 304, 1781-1781.	6.0	229
107	Genetic Polymorphism of 11 Allozyme Loci in Populations of Wall Lizards (<i>Podarcis</i> sp.) from the Iberian Peninsula and North Africa. <i>Biochemical Genetics</i> , 2003, 41, 343-359.	0.8	24
108	Biochemical and Population Genetics of the Rabbit, <i>Oryctolagus cuniculus</i> , Carbonic Anhydrases I and II, from the Iberian Peninsula and France. <i>Biochemical Genetics</i> , 2003, 41, 391-404.	0.8	9

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109	Ancient introgression of <i>Lepus timidus</i> mtDNA into <i>L. granatensis</i> and <i>L. europaeus</i> in the Iberian Peninsula. <i>Molecular Phylogenetics and Evolution</i> , 2003, 27, 70-80.	1.2	112
110	Genetic structure of eighteen local south European beef cattle breeds by comparative F-statistics analysis. <i>Journal of Animal Breeding and Genetics</i> , 2003, 120, 73-87.	0.8	46
111	Gene-culture coevolution between cattle milk protein genes and human lactase genes. <i>Nature Genetics</i> , 2003, 35, 311-313.	9.4	371
112	Genetic Characterization of Southwestern European Bovine Breeds: A Historical and Biogeographical Reassessment With a Set of 16 Microsatellites. , 2003, 94, 243-250.		78
113	Complex patterns of genetic diversity within <i>Lacerta (Teira) perspicillata</i> : Preliminary evidence from 12S rRNA sequence data. <i>Amphibia - Reptilia</i> , 2003, 24, 386-390.	0.1	13
114	Reproductive cycle of the golden-striped salamander <i>Chioglossa lusitanica</i> (Caudata, Salamandridae) in NW Portugal. <i>Amphibia - Reptilia</i> , 2003, 24, 1-12.	0.1	9
115	POSTGLACIAL DISPERSAL OF THE EUROPEAN RABBIT (<i>ORYCTOLAGUS CUNICULUS</i>) ON THE IBERIAN PENINSULA RECONSTRUCTED FROM NESTED CLADE AND MISMATCH ANALYSES OF MITOCHONDRIAL DNA GENETIC VARIATION. <i>Evolution; International Journal of Organic Evolution</i> , 2002, 56, 792.	1.1	4
116	Microsatellite Variation and Evolution of the Human Duffy Blood Group Polymorphism. <i>Molecular Biology and Evolution</i> , 2002, 19, 1802-1806.	3.5	24
117	Hotspot variation at the CH2-CH3 interface of leporid IgG antibodies (<i>Oryctolagus, Sylvilagus</i> and) Tj ETQq1 1 0.784314 rgBT /Overlo	1.2	10
118	Complex biogeographical distribution of genetic variation within <i>Podarcis wall</i> lizards across the Strait of Gibraltar. <i>Journal of Biogeography</i> , 2002, 29, 1257-1262.	1.4	93
119	Evidence for a geographical cline of casein haplotypes in Portuguese cattle breeds. <i>Animal Genetics</i> , 2002, 33, 295-300.	0.6	29
120	Restriction fragment alleles of the rabbitIGHGgenes with reference to the rabbitIGHGCH2or e locus polymorphism. <i>Animal Genetics</i> , 2002, 33, 309-311.	0.6	8
121	POSTGLACIAL DISPERSAL OF THE EUROPEAN RABBIT (<i>ORYCTOLAGUS CUNICULUS</i>) ON THE IBERIAN PENINSULA RECONSTRUCTED FROM NESTED CLADE AND MISMATCH ANALYSES OF MITOCHONDRIAL DNA GENETIC VARIATION. <i>Evolution; International Journal of Organic Evolution</i> , 2002, 56, 792-803.	1.1	100
122	Genetic polymorphism of antithrombin III, haptoglobin, and haemopexin in wild and domestic European rabbits. <i>Biochemical Genetics</i> , 2002, 40, 387-393.	0.8	5
123	Genetic polymorphism of the 17th exon at porcine RYR1 locus: a new variant in a local Portuguese pig breed demonstrated by SSCP analysis. <i>Journal of Animal Breeding and Genetics</i> , 2001, 118, 271-274.	0.8	2
124	Genetic analysis and mapping of biochemical markers in an F2 intercross of two inbred strains of the rabbit (<i>Oryctolagus cuniculus</i>). <i>Biochemical Genetics</i> , 2001, 39, 169-178.	0.8	8
125	Stationary Distributions of Microsatellite Loci Between Divergent Population Groups of the European Rabbit (<i>Oryctolagus cuniculus</i>). <i>Molecular Biology and Evolution</i> , 2001, 18, 2169-2178.	3.5	83
126	Age structure and growth pattern in two populations of the golden-striped salamander <i>Chioglossa lusitanica</i> (Caudata, Salamandridae). <i>Amphibia - Reptilia</i> , 2001, 22, 55-68.	0.1	47

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127	Phylogeography of the European rabbit (<i>Oryctolagus cuniculus</i>) in the Iberian Peninsula inferred from RFLP analysis of the cytochrome b gene. <i>Heredity</i> , 2000, 85, 307-317.	1.2	116
128	Genetic subdivision, glacial refugia and postglacial recolonization in the golden-striped salamander, <i>Chioglossa lusitanica</i> (Amphibia: Urodela). <i>Molecular Ecology</i> , 2000, 9, 771-781.	2.0	102
129	Absence of a genetic bottleneck in a wild rabbit (<i>Oryctolagus cuniculus</i>) population exposed to a severe viral epizootic. <i>Molecular Ecology</i> , 2000, 9, 1253-1264.	2.0	44
130	New genetic variation in European hares, <i>Lepus granatensis</i> and <i>L. europaeus</i> . <i>Biochemical Genetics</i> , 2000, 38, 87-96.	0.8	15
131	Extensive genetic polymorphism of peptidases A, B, C, and D, in wild rabbit (<i>Oryctolagus cuniculus</i>) populations from the Iberian Peninsula. <i>Biochemical Genetics</i> , 1999, 37, 237-249.	0.8	11
132	Cytonuclear disequilibria in wild populations of rabbit (<i>Oryctolagus cuniculus</i> L.) suggest unequal allele turnover rates at the b locus (IGKC1). <i>Immunogenetics</i> , 1999, 49, 629-643.	1.2	23
133	Genetic polymorphism of properdin factor B (BF) in domestic rabbit. <i>Animal Genetics</i> , 1998, 29, 135-137.	0.6	8
134	Genetic polymorphism of rabbit (<i>Oryctolagus cuniculus</i>) tissue acid phosphatases (ACP2 and ACP3). <i>Comparative Biochemistry and Physiology - B Biochemistry and Molecular Biology</i> , 1998, 120, 405-409.	0.7	13
135	Genetic variation in some populations of the golden-striped salamander, <i>Chioglossa lusitanica</i> (Amphibia: Urodela), in Portugal. <i>Biochemical Genetics</i> , 1997, 35, 371-381.	0.8	10
136	Genetic polymorphism of adenosine deaminase (ADA; E.C. 3.5.4.4.) in allis shad, <i>Alosa alosa</i> and twaite shad, <i>Alosa fallax</i> . <i>Journal of Fish Biology</i> , 1993, 43, 951-953.	0.7	1
137	Separation of human alloalbumin variants by isoelectric focusing. <i>Electrophoresis</i> , 1991, 12, 313-314.	1.3	5
138	Genetic polymorphism of α -aminolaevulinic acid dehydratase (E.C. 4.2.1.24, ALAD) in the domestic rabbit. <i>Animal Genetics</i> , 1990, 21, 217-219.	0.6	8
139	Biochemical and genetic studies on rabbit hemoglobin. II. Electrophoretic polymorphism of the β -chain. <i>Biochemical Genetics</i> , 1990, 28, 117-122.	0.8	16
140	Biochemical and genetic studies on rabbit hemoglobin. I. Electrophoretic polymorphism of the β^2 chain. <i>Biochemical Genetics</i> , 1989, 27, 673-678.	0.8	16
141	Documenting the advantages and limitations of different classes of molecular markers in a well-established phylogeographic context: lessons from the Iberian endemic Golden-striped salamander, <i>Chioglossa lusitanica</i> (Caudata: Salamandridae). <i>Biological Journal of the Linnean Society</i> , 0, 95, 371-387.	0.7	25
142	Nested clade analysis and the genetic evidence for population expansion in the phylogeography of the golden-striped salamander, <i>Chioglossa lusitanica</i> (Amphibia: Urodela). , 0, .		3