

# Paulo CÃ©lio Alves

## List of Publications by Year in descending order

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Version: 2024-02-01

172  
papers

6,573  
citations

66234

42  
h-index

91712

69  
g-index

177  
all docs

177  
docs citations

177  
times ranked

7461  
citing authors

#	ARTICLE	IF	CITATIONS
1	Genomics and the challenging translation into conservation practice. Trends in Ecology and Evolution, 2015, 30, 78-87.	4.2	469
2	Advancing ecological understandings through technological transformations in noninvasive genetics. Molecular Ecology Resources, 2009, 9, 1279-1301.	2.2	296
3	Adaptive introgression underlies polymorphic seasonal camouflage in snowshoe hares. Science, 2018, 360, 1355-1358.	6.0	234
4	Invasion from the cold past: extensive introgression of mountain hare ( <i>Lepus timidus</i> ) mitochondrial DNA into three other hare species in northern Iberia. Molecular Ecology, 2005, 14, 2459-2464.	2.0	183
5	Plasticity in circadian activity patterns of mesocarnivores in Southwestern Europe: implications for species coexistence. Behavioral Ecology and Sociobiology, 2014, 68, 1403-1417.	0.6	183
6	Reference-Free Population Genomics from Next-Generation Transcriptome Data and the Vertebrate–Invertebrate Gap. PLoS Genetics, 2013, 9, e1003457.	1.5	157
7	Proposal for a unified classification system and nomenclature of lagoviruses. Journal of General Virology, 2017, 98, 1658-1666.	1.3	148
8	Catch Me If You Can: Diel Activity Patterns of Mammalian Prey and Predators. Ethology, 2013, 119, 1044-1056.	0.5	128
9	Ancient introgression of <i>Lepus timidus</i> mtDNA into <i>L. granatensis</i> and <i>L. europaeus</i> in the Iberian Peninsula. Molecular Phylogenetics and Evolution, 2003, 27, 70-80.	1.2	112
10	The ubiquitous mountain hare mitochondria: multiple introgressive hybridization in hares, genus <i>Lepus</i> . Philosophical Transactions of the Royal Society B: Biological Sciences, 2008, 363, 2831-2839.	1.8	111
11	Recurrent Introgression of Mitochondrial DNA Among Hares ( <i>Lepus</i> spp.) Revealed by Species-Tree Inference and Coalescent Simulations. Systematic Biology, 2012, 61, 367.	2.7	111
12	Function and underlying mechanisms of seasonal colour moulting in mammals and birds: what keeps them changing in a warming world?. Biological Reviews, 2018, 93, 1478-1498.	4.7	109
13	Molecular analysis of hybridisation between wild and domestic cats ( <i>Felis silvestris</i> ) in Portugal: implications for conservation. Conservation Genetics, 2008, 9, 1-11.	0.8	100
14	Adapted conservation measures are required to save the Iberian lynx in a changing climate. Nature Climate Change, 2013, 3, 899-903.	8.1	96
15	The rise and fall of the mountain hare ( <i>Lepus timidus</i> ) during Pleistocene glaciations: expansion and retreat with hybridization in the Iberian Peninsula. Molecular Ecology, 2006, 16, 605-618.	2.0	95
16	Hybridization versus conservation: are domestic cats threatening the genetic integrity of wildcats ( <i>Felis silvestris silvestris</i> ) in Iberian Peninsula?. Philosophical Transactions of the Royal Society B: Biological Sciences, 2008, 363, 2953-2961.	1.8	91
17	Winter color polymorphisms identify global hot spots for evolutionary rescue from climate change. Science, 2018, 359, 1033-1036.	6.0	91
18	Spatial ecology of the European wildcat in a Mediterranean ecosystem: dealing with small radio-tracking datasets in species conservation. Journal of Zoology, 2009, 279, 27-35.	0.8	89

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19	New Variant of Rabbit Hemorrhagic Disease Virus, Portugal, 2012–2013. <i>Emerging Infectious Diseases</i> , 2013, 19, 1900-2.	2.0	86
20	Phylogeography of roe deer ( <i>Capreolus capreolus</i> ) populations: the effects of historical genetic subdivisions and recent nonequilibrium dynamics. <i>Molecular Ecology</i> , 2004, 13, 3071-3083.	2.0	80
21	The Elusive Nature of Adaptive Mitochondrial DNA Evolution of an Arctic Lineage Prone to Frequent Introgression. <i>Genome Biology and Evolution</i> , 2014, 6, 886-896.	1.1	78
22	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
23	Is the New Variant RHDV Replacing Genogroup 1 in Portuguese Wild Rabbit Populations?. <i>Viruses</i> , 2015, 7, 27-36.	1.5	66
24	Applying genomic data in wildlife monitoring: Development guidelines for genotyping degraded samples with reduced single nucleotide polymorphism panels. <i>Molecular Ecology Resources</i> , 2020, 20, 662-680.	2.2	64
25	Phylogeography of the brown hare ( <i>Lepus europaeus</i> ) in Europe: a legacy of south-eastern Mediterranean refugia?. <i>Journal of Biogeography</i> , 2009, 36, 515-528.	1.4	63
26	Integrative approaches to guide conservation decisions: Using genomics to define conservation units and functional corridors. <i>Molecular Ecology</i> , 2018, 27, 3452-3465.	2.0	63
27	Effect of microsatellite selection on individual and population genetic inferences: an empirical study using cross-specific and species-specific amplifications. <i>Molecular Ecology Resources</i> , 2015, 15, 747-760.	2.2	61
28	Ecological traits and the spatial structure of competitive coexistence among carnivores. <i>Ecology</i> , 2020, 101, e03059.	1.5	61
29	Feeding ecological knowledge: the underutilised power of faecal <i>scp</i> DNA approaches for carnivore diet analysis. <i>Mammal Review</i> , 2019, 49, 97-112.	2.2	60
30	Cryptic speciation in the field vole: a multilocus approach confirms three highly divergent lineages in <i>scp</i> Eurasia. <i>Molecular Ecology</i> , 2012, 21, 6015-6032.	2.0	59
31	Genetic structure of wildcat ( <i>Felis silvestris</i> ) populations in Italy. <i>Ecology and Evolution</i> , 2013, 3, 2443-2458.	0.8	58
32	Disease-mediated bottom-up regulation: An emergent virus affects a keystone prey, and alters the dynamics of trophic webs. <i>Scientific Reports</i> , 2016, 6, 36072.	1.6	58
33	Mitochondrial phylogeography of the European wild boar: the effect of climate on genetic diversity and spatial lineage sorting across Europe. <i>Journal of Biogeography</i> , 2014, 41, 987-998.	1.4	56
34	Seasonal variation in the reproductive activity of the wild rabbit ( <i>Oryctolagus cuniculus algirus</i> ) in a Mediterranean ecosystem. <i>Wildlife Research</i> , 2002, 29, 165.	0.7	55
35	Influence of habitat management on the abundance and diet of wild rabbit ( <i>Oryctolagus cuniculus</i> ) Tj ETQq1 1 0.784314 rgBT /Overlock 487-496.	0.7	55
36	Genetic identification of <i>scp</i> Iberian rodent species using both mitochondrial and nuclear loci: application to noninvasive sampling. <i>Molecular Ecology Resources</i> , 2013, 13, 43-56.	2.2	55

#	ARTICLE	IF	CITATIONS
37	Insights into the evolution of the new variant rabbit haemorrhagic disease virus (Gl.2) and the identification of novel recombinant strains. <i>Transboundary and Emerging Diseases</i> , 2018, 65, 983-992.	1.3	52
38	Toward a genome-wide approach for detecting hybrids: informative SNPs to detect introgression between domestic cats and European wildcats ( <i>Felis silvestris</i> ). <i>Heredity</i> , 2015, 115, 195-205.	1.2	51
39	Evolution of rabbit haemorrhagic disease virus (RHDV) in the European rabbit ( <i>Oryctolagus</i> ) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50	0.8	50
40	Introgression of mitochondrial DNA among <i>Myodes</i> voles: consequences for energetics?. <i>BMC Evolutionary Biology</i> , 2011, 11, 355.	3.2	50
41	Parapatric species and the implications for climate change studies: a case study on hares in Europe. <i>Global Change Biology</i> , 2012, 18, 1509-1519.	4.2	49
42	European wildcat populations are subdivided into five main biogeographic groups: consequences of Pleistocene climate changes or recent anthropogenic fragmentation?. <i>Ecology and Evolution</i> , 2016, 6, 3-22.	0.8	49
43	Factors affecting the (in)accuracy of mammalian mesocarnivore scat identification in South-western Europe. <i>Journal of Zoology</i> , 2013, 289, 243-250.	0.8	48
44	The Legacy of Recurrent Introgression during the Radiation of Hares. <i>Systematic Biology</i> , 2021, 70, 593-607.	2.7	47
45	Evaluation of attractants for non-invasive studies of Iberian carnivore communities. <i>Wildlife Research</i> , 2011, 38, 446.	0.7	45
46	Genetic non-invasive sampling (gNIS) as a cost-effective tool for monitoring elusive small mammals. <i>European Journal of Wildlife Research</i> , 2018, 64, 1.	0.7	45
47	Have the cake and eat it: Optimizing nondestructive DNA metabarcoding of macroinvertebrate samples for freshwater biomonitoring. <i>Molecular Ecology Resources</i> , 2019, 19, 863-876.	2.2	44
48	Population genetics of cape and brown hares ( <i>Lepus capensis</i> and <i>L. europaeus</i> ): A test of Petter's hypothesis of conspecificity. <i>Biochemical Systematics and Ecology</i> , 2008, 36, 22-39.	0.6	43
49	Genetic diversity of wild boar populations and domestic pig breeds ( <i>Sus scrofa</i> ) in South-western Europe. <i>Biological Journal of the Linnean Society</i> , 2010, 101, 797-822.	0.7	42
50	Epidemiology of RHDV2 ( <i>Lagovirus europaeus</i> /Gl.2) in free-living wild European rabbits in Portugal. <i>Transboundary and Emerging Diseases</i> , 2018, 65, e373-e382.	1.3	41
51	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
52	The hidden history of the snowshoe hare, <i>Lepus americanus</i> : extensive mitochondrial DNA introgression inferred from multilocus genetic variation. <i>Molecular Ecology</i> , 2014, 23, 4617-4630.	2.0	40
53	Evidence for genetic similarity of two allopatric European hares ( <i>Lepus corsicanus</i> and <i>L.</i> ) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 1191-1197.	1.2	39
54	Molecular and ecological signs of mitochondrial adaptation: consequences for introgression?. <i>Heredity</i> , 2014, 113, 277-286.	1.2	37

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55	Species identification using a small nuclear gene fragment: application to sympatric wild carnivores from South-western Europe. <i>Conservation Genetics</i> , 2010, 11, 1023-1032.	0.8	36
56	Range expansion underlies historical introgressive hybridization in the Iberian hare. <i>Scientific Reports</i> , 2017, 7, 40788.	1.6	35
57	Genomic approaches to identify hybrids and estimate admixture times in European wildcat populations. <i>Scientific Reports</i> , 2019, 9, 11612.	1.6	34
58	Conservation implications of the evolutionary history and genetic diversity hotspots of the snowshoe hare. <i>Molecular Ecology</i> , 2014, 23, 2929-2942.	2.0	32
59	Molecular bases of genetic diversity and evolution of the immunoglobulin heavy chain variable region (IGHV) gene locus in leporids. <i>Immunogenetics</i> , 2011, 63, 397-408.	1.2	31
60	Past, Present and Future Distributions of an Iberian Endemic, <i>Lepus granatensis</i> : Ecological and Evolutionary Clues from Species Distribution Models. <i>PLoS ONE</i> , 2012, 7, e51529.	1.1	31
61	Range-wide patterns of human-mediated hybridisation in European wildcats. <i>Conservation Genetics</i> , 2020, 21, 247-260.	0.8	31
62	Home-loving boreal hare mitochondria survived several invasions in Iberia: the relative roles of recurrent hybridisation and allele surfing. <i>Heredity</i> , 2014, 112, 265-273.	1.2	30
63	Precision Medicine in Cats: Novel Niemann-Pick Type C1 Diagnosed by Whole-Genome Sequencing. <i>Journal of Veterinary Internal Medicine</i> , 2017, 31, 539-544.	0.6	30
64	A genomic map of clinal variation across the European rabbit hybrid zone. <i>Molecular Ecology</i> , 2018, 27, 1457-1478.	2.0	30
65	Endemic Sand Dune Vegetation of the Northwest Iberian Peninsula: Diversity, Dynamics, and Significance for Bioindication and Monitoring of Coastal Landscapes. <i>Journal of Coastal Research</i> , 2008, 2, 113-121.	0.1	29
66	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
67	Efficiency of hair snares and camera traps to survey mesocarnivore populations. <i>European Journal of Wildlife Research</i> , 2014, 60, 279-289.	0.7	29
68	Females know better: Sex-biased habitat selection by the European wildcat. <i>Ecology and Evolution</i> , 2018, 8, 9464-9477.	0.8	29
69	Red deer in Iberia: Molecular ecological studies in a southern refugium and inferences on European postglacial colonization history. <i>PLoS ONE</i> , 2019, 14, e0210282.	1.1	29
70	Mutations in the Kinesin-2 Motor KIF3B Cause an Autosomal-Dominant Ciliopathy. <i>American Journal of Human Genetics</i> , 2020, 106, 893-904.	2.6	29
71	Quantification of the Animal Tuberculosis Multi-Host Community Offers Insights for Control. <i>Pathogens</i> , 2020, 9, 421.	1.2	29
72	Genetic diversity within scorpions of the genus <i>Buthus</i> from the Iberian Peninsula: mitochondrial DNA sequence data indicate additional distinct cryptic lineages. <i>Journal of Arachnology</i> , 2010, 38, 206-211.	0.3	28

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73	Niche partitioning at the edge of the range: a multidimensional analysis with sympatric martens. Journal of Mammalogy, 2016, 97, 928-939.	0.6	28
74	The Microtus voles: Resolving the phylogeny of one of the most speciose mammalian genera using genomics. Molecular Phylogenetics and Evolution, 2018, 125, 85-92.	1.2	28
75	Stepping up from wildlife disease surveillance to integrated wildlife monitoring in Europe. Research in Veterinary Science, 2022, 144, 149-156.	0.9	28
76	Genetic Diversity of Maghrebian <i>Hottentotta</i> (Scorpiones: Buthidae) Scorpions Based on CO1: New Insights on the Genus Phylogeny and Distribution. African Invertebrates, 2011, 52, 135-143.	0.5	27
77	The transcriptional landscape of seasonal coat colour moult in the snowshoe hare. Molecular Ecology, 2017, 26, 4173-4185.	2.0	27
78	Reproductive biology of the Iberian hare, <i>Lepus granatensis</i> , in Portugal. Mammalian Biology, 2002, 67, 358-371.	0.8	26
79	Endemic species may have complex histories: within-refugium phylogeography of an endangered Iberian vole. Molecular Ecology, 2017, 26, 951-967.	2.0	26
80	Genetic identification of endangered North African ungulates using noninvasive sampling. Molecular Ecology Resources, 2015, 15, 652-661.	2.2	25
81	Field experimental vaccination campaigns against myxomatosis and their effectiveness in the wild. Vaccine, 2009, 27, 6998-7002.	1.7	24
82	Detection of RHDV strains in the Iberian hare ( <i>Lepus granatensis</i> ): earliest evidence of rabbit lagovirus cross-species infection. Veterinary Research, 2014, 45, 94.	1.1	24
83	Reply to Garner et al.. Trends in Ecology and Evolution, 2016, 31, 83-84.	4.2	24
84	Detection of RHDV strains in the Iberian hare ( <i>Lepus granatensis</i> ): earliest evidence of rabbit lagovirus cross-species infection. Veterinary Research, 2014, 45, 94.	1.1	24
85	The impact of management practices and past demographic history on the genetic diversity of red deer ( <i>Cervus elaphus</i> ): an assessment of population and individual fitness. Biological Journal of the Linnean Society, 2014, 111, 209-223.	0.7	23
86	Early-Onset Progressive Retinal Atrophy Associated with an IQCB1 Variant in African Black-Footed Cats ( <i>Felis nigripes</i> ). Scientific Reports, 2017, 7, 43918.	1.6	22
87	Evolutionary relationships among hares from North Africa ( <i>Lepus</i> sp. or <i>Lepus</i> spp.), cape hares ( <i>L.</i> ) <i>Tj ETQq1 1 0.784314 rgBT /Overlock</i> allozyme data. Journal of Zoological Systematics and Evolutionary Research, 2006, 44, 88-99.	0.6	21
88	Diet of the Iberian hare ( <i>Lepus granatensis</i> ) in a mountain ecosystem. European Journal of Wildlife Research, 2008, 54, 571-579.	0.7	21
89	The usefulness of field data and hunting statistics in the assessment of wild rabbit ( <i>Oryctolagus</i> ) <i>Tj ETQq1 1 0.784314 rgBT /Overlock</i>	0.7	21
90	Range dynamics driven by Quaternary climate oscillations explain the distribution of introgressed mtDNA of <i>Lepus timidus</i> origin in hares from the Iberian Peninsula. Journal of Biogeography, 2015, 42, 1727-1735.	1.4	21

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91	The effects of a motorway on movement behaviour and gene flow in a forest carnivore: Joint evidence from road mortality, radio tracking and genetics. <i>Landscape and Urban Planning</i> , 2018, 178, 217-227.	3.4	20
92	Werewolf, There Wolf: Variants in Hairless Associated with Hypotrichia and Roaning in the Lykoi Cat Breed. <i>Genes</i> , 2020, 11, 682.	1.0	20
93	Hares in Corsica: high prevalence of <i>Lepus corsicanus</i> and hybridization with introduced <i>L. europaeus</i> and <i>L. granatensis</i> . <i>European Journal of Wildlife Research</i> , 2011, 57, 313-321.	0.7	19
94	Molecular phylogeny of the Western Palaearctic <i>Cordulegaster</i> taxa (Odonata: Anisoptera: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50	0.7	19
95	A Critically Endangered new dragonfly species from Morocco: <i>Onychogomphus boudoti</i> sp. nov. (Odonata: Gomphidae). <i>Zootaxa</i> , 2014, 3856, 349-65.	0.2	19
96	Can we predict habitat quality from space? A multi-indicator assessment based on an automated knowledge-driven system. <i>International Journal of Applied Earth Observation and Geoinformation</i> , 2015, 37, 106-113.	1.4	19
97	Tuberculosis, genetic diversity and fitness in the red deer, <i>Cervus elaphus</i> . <i>Infection, Genetics and Evolution</i> , 2016, 43, 203-212.	1.0	19
98	LaGomiCsâ€™ Lagomorph Genomics Consortium: An International Collaborative Effort for Sequencing the Genomes of an Entire Mammalian Order. <i>Journal of Heredity</i> , 2016, 107, 295-308.	1.0	19
99	Genetic diversity within <i>Scorpio maurus</i> (Scorpiones: Scorpionidae) from morocco: Preliminary evidence based on CO1 mitochondrial DNA sequences. <i>Biologia (Poland)</i> , 2008, 63, 1157-1160.	0.8	18
100	Sequencing of modern <i>Lepus</i> VDJ genes shows that the usage of VHn genes has been retained in both <i>Oryctolagus</i> and <i>Lepus</i> that diverged 12 million years ago. <i>Immunogenetics</i> , 2013, 65, 777-784.	1.2	18
101	Leporid immunoglobulin G shows evidence of strong selective pressure on the hinge and CH3 domains. <i>Open Biology</i> , 2014, 4, 140088.	1.5	18
102	Biometrical analysis reveals major differences between the two subspecies of the European rabbit. <i>Biological Journal of the Linnean Society</i> , 2015, 116, 106-116.	0.7	18
103	Spatial climate dynamics in the Iberian Peninsula since 15â€™000â€™yrâ€™BP. <i>Climate of the Past</i> , 2016, 12, 1137-1149.	1.4	18
104	Optimizing cameraâ€™trapping protocols for characterizing mesocarnivore communities in southâ€™western Europe. <i>Journal of Zoology</i> , 2017, 301, 23-31.	0.8	18
105	Evidence for niche similarities in the allopatric sister species <i>Lepus castroviejoi</i> and <i>Lepus corsicanus</i> . <i>Journal of Biogeography</i> , 2014, 41, 977-986.	1.4	17
106	A FAS-ligand variant associated with autoimmune lymphoproliferative syndrome in cats. <i>Mammalian Genome</i> , 2017, 28, 47-55.	1.0	17
107	Combining distribution modelling and non-invasive genetics to improve range shift forecasting. <i>Ecological Modelling</i> , 2015, 297, 171-179.	1.2	16
108	First genome-wide CNV mapping in <i>FELIS CATUS</i> using next generation sequencing data. <i>BMC Genomics</i> , 2018, 19, 895.	1.2	16



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109	Transcriptomic regulation of seasonal coat color change in hares. <i>Ecology and Evolution</i> , 2020, 10, 1180-1192.	0.8	16
110	Evolutionary history of two cryptic species of northern African jerboas. <i>BMC Evolutionary Biology</i> , 2020, 20, 26.	3.2	16
111	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
112	Urban Habitats Biodiversity Assessment (UrHBA): a standardized procedure for recording biodiversity and its spatial distribution in urban environments. <i>Landscape Ecology</i> , 2017, 32, 1753-1770.	1.9	15
113	Genome-wide associations identify novel candidate loci associated with genetic susceptibility to tuberculosis in wild boar. <i>Scientific Reports</i> , 2018, 8, 1980.	1.6	15
114	Recent range contractions in the globally threatened Pyrenean desman highlight the importance of stream headwater refugia. <i>Animal Conservation</i> , 2018, 21, 515-525.	1.5	15
115	Candidate genes underlying heritable differences in reproductive seasonality between wild and domestic rabbits. <i>Animal Genetics</i> , 2015, 46, 418-425.	0.6	14
116	Characterization of old RHDV strains by complete genome sequencing identifies a novel genetic group. <i>Scientific Reports</i> , 2017, 7, 13599.	1.6	14
117	Combining genetic non-invasive sampling with spatially explicit capture-recapture models for density estimation of a patchily distributed small mammal. <i>European Journal of Wildlife Research</i> , 2018, 64, 1.	0.7	14
118	Molecular and morphological insights into the origin of the invasive greater white-toothed shrew ( <i>Crocidura russula</i> ) in Ireland. <i>Biological Invasions</i> , 2016, 18, 857-871.	1.2	13
119	An Annotated Draft Genome of the Mountain Hare ( <i>Lepus timidus</i> ). <i>Genome Biology and Evolution</i> , 2020, 12, 3656-3662.	1.1	13
120	Evidence of autumn reproduction in female European hares ( <i>Lepus europaeus</i> ) from southern Europe. <i>European Journal of Wildlife Research</i> , 2008, 54, 581-587.	0.7	12
121	Estimating home-range size: when to include a third dimension?. <i>Ecology and Evolution</i> , 2013, 3, 2285-2295.	0.8	12
122	Genetic distinctiveness of the damselfly <i>Coenagrion puella</i> in North Africa: an overlooked and endangered taxon. <i>Conservation Genetics</i> , 2016, 17, 985-991.	0.8	12
123	The evolutionary history of the Cape hare ( <i>Lepus capensis sensu lato</i> ): insights for systematics and biogeography. <i>Heredity</i> , 2019, 123, 634-646.	1.2	12
124	Comparative Proteomics Identifies Host Immune System Proteins Affected by Infection with <i>Mycobacterium bovis</i> . <i>PLoS Neglected Tropical Diseases</i> , 2016, 10, e0004541.	1.3	12
125	Environmental factors have little influence on the reproductive activity of the Iberian hare ( <i>Lepus</i> )	0.7	11
126	Mountain hare transcriptome and diagnostic markers as resources to monitor hybridization with European hares. <i>Scientific Data</i> , 2017, 4, 170178.	2.4	11



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127	Deciphering Anthropogenic Effects on the Genetic Background of the Red Deer in the Iberian Peninsula. <i>Frontiers in Ecology and Evolution</i> , 2020, 8, .	1.1	11
128	Hotspot variation at the CH2-CH3 interface of leporid IgG antibodies ( <i>Oryctolagus</i> , <i>Sylvilagus</i> and) Tj ETQq0 0 0 rgBT/Overlook 10 Tf 50	1.2	10
129	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
130	Local extinctions and range contraction of the endangered <i>Coenagrion mercuriale</i> in North Africa. <i>International Journal of Odonatology</i> , 2015, 18, 137-152.	0.5	9
131	Feline mitochondrial DNA sampling for forensic analysis: When enough is enough!. <i>Forensic Science International: Genetics</i> , 2015, 16, 52-57.	1.6	9
132	Ecotypes and evolutionary significant units in endangered North African gazelles. <i>Biological Journal of the Linnean Society</i> , 2017, 122, 286-300.	0.7	9
133	Mining the 99 Lives Cat Genome Sequencing Consortium database implicates genes and variants for the <i>Ticked</i> locus in domestic cats ( <i>Felis catus</i> ). <i>Animal Genetics</i> , 2021, 52, 321-332.	0.6	9
134	Restriction fragment alleles of the rabbit <i>IGHG</i> genes with reference to the rabbit <i>IGHG</i> <i>CH2</i> or <i>e</i> locus polymorphism. <i>Animal Genetics</i> , 2002, 33, 309-311.	0.6	8
135	Positive selection on the mitochondrial <i>ATP synthase 6</i> and the <i>NADH dehydrogenase 2</i> genes across 22 hare species (genus <i>Lepus</i> ). <i>Journal of Zoological Systematics and Evolutionary Research</i> , 2018, 56, 428-443.	0.6	8
136	Gastrointestinal parasite infestation in the alpine mountain hare ( <i>Lepus timidus varronis</i> ): Are abiotic environmental factors such as elevation, temperature and precipitation affecting prevalence of parasite species?. <i>International Journal for Parasitology: Parasites and Wildlife</i> , 2019, 9, 202-208.	0.6	8
137	Drivers of survival in a small mammal of conservation concern: An assessment using extensive genetic non-invasive sampling in fragmented farmland. <i>Biological Conservation</i> , 2019, 230, 131-140.	1.9	8
138	Overview of Lagomorph Research: What we have learned and what we still need to do. , 2008, , 381-391.		8
139	The evolutionary pathways for local adaptation in mountain hares. <i>Molecular Ecology</i> , 2022, 31, 1487-1503.	2.0	8
140	Patterns of genetic diversity within and between <i>Myotis d. daubentonii</i> and <i>M. d. nathalinae</i> derived from cytochrome b mtDNA sequence data. <i>Acta Chiropterologica</i> , 2007, 9, 379-389.	0.2	7
141	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
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