Paulo CÃ©lio Alves

List of Publications by Year
in descending order

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	Stepping up from wildlife disease surveillance to integrated wildlife monitoring in Europe. Research in Veterinary Science, 2022, 144, 149-156.	0.9	28
2	The evolutionary pathways for local adaptation in mountain hares. Molecular Ecology, 2022, 31, 1487-1503.	2.0	8
3	MAMMALS IN PORTUGAL : A data set of terrestrial, volant, and marine mammal occurrences in Portugal. Ecology, 2022, , e3654.	1.5	1
4	Genetic integrity of European wildcats: Variation across biomes mandates geographically tailored conservation strategies. Biological Conservation, 2022, 268, 109518.	1.9	4
5	Comfort over safety: thermoregulation overshadows predation risk effects in the activity of a keystone prey. Journal of Zoology, 2022, 316, 209-222.	0.8	7
6	Bagaza Virus in Wild Birds, Portugal, 2021. Emerging Infectious Diseases, 2022, 28, 1504-1506.	2.0	4
7	Multiâ€event captureâ€“recapture models estimate the diagnostic performance of serological tests for myxoma and rabbit haemorrhagic disease viruses in the absence of reference samples. Transboundary and Emerging Diseases, 2022, 69, .	1.3	3
8	The Legacy of Recurrent Introgression during the Radiation of Hares. Systematic Biology, 2021, 70, 593-607.	2.7	47
9	Mining the 99 Lives Cat Genome Sequencing Consortium database implicates genes and variants for the Ticked locus in domestic cats (FelisÂcatus). Animal Genetics, 2021, 52, 321-332.	0.6	9
10	Multi-omic analyses in Abyssinian cats with primary renal amyloid deposits. Scientific Reports, 2021, 11, 8339.	1.6	6
11	Iberian hares with anciently introgressed mitochondrial DNA express a marginal environmental niche. Journal of Biogeography, 2021, 48, 2328-2336.	1.4	6
12	Assessing changes in stream macroinvertebrate communities across ecological gradients using morphological versus DNA metabarcoding approaches. Science of the Total Environment, 2021, 797, 149030.	3.9	3
13	Sex and Age-Specific Hematology and Biochemistry Reference Intervals of Live Iberian Hares (Lepus) Tj ETQq1 1 0.784314 rgBT /Overl	0.3	2
14	Integrating multiple datasets into spatially-explicit capture-recapture models to estimate the abundance of a locally scarce felid. Biodiversity and Conservation, 2021, 30, 4317-4335.	1.2	3
15	European Rabbit Oryctolagus cuniculus (Linnaeus, 1758). Handbook of the Mammals of Europe, 2021, , 1-39.	0.1	2
16	An Annotated Draft Genome of the Mountain Hare (Lepus timidus). Genome Biology and Evolution, 2020, 12, 3656-3662.	1.1	13
17	Effect of landscape type, elevation, vegetation period, and taxonomic plant identification level on diet preferences of Alpine mountain hares (Lepus timidus varronis). European Journal of Wildlife Research, 2020, 66, 1.	0.7	6
18	Mutations in the Kinesin-2 Motor KIF3B Cause an Autosomal-Dominant Ciliopathy. American Journal of Human Genetics, 2020, 106, 893-904.	2.6	29

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19	A Deletion in GDF7 is Associated with a Heritable Forebrain Commissural Malformation Concurrent with Ventriculomegaly and Interhemispheric Cysts in Cats. Genes, 2020, 11, 672.	1.0	7
20	Quantification of the Animal Tuberculosis Multi-Host Community Offers Insights for Control. Pathogens, 2020, 9, 421.	1.2	29
21	Werewolf, There Wolf: Variants in Hairless Associated with Hypotrichia and Roaning in the Lykoi Cat Breed. Genes, 2020, 11, 682.	1.0	20
22	Transcriptomic regulation of seasonal coat color change in hares. Ecology and Evolution, 2020, 10, 1180-1192.	0.8	16
23	Taxonomic identification of Madagascarâ€™s free-ranging â€œforest catsâ€• Conservation Genetics, 2020, 21, 443-451.	0.8	5
24	Evolutionary history of two cryptic species of northern African jerboas. BMC Evolutionary Biology, 2020, 20, 26.	3.2	16
25	Genetic diversity in natural range remnants of the critically endangered hirola antelope. Zoological Journal of the Linnean Society, 2020, 190, 384-395.	1.0	3
26	Range-wide patterns of human-mediated hybridisation in European wildcats. Conservation Genetics, 2020, 21, 247-260.	0.8	31
27	Applying genomic data in wildlife monitoring: Development guidelines for genotyping degraded samples with reduced single nucleotide polymorphism panels. Molecular Ecology Resources, 2020, 20, 662-680.	2.2	64
28	Deciphering Anthropogenic Effects on the Genetic Background of the Red Deer in the Iberian Peninsula. Frontiers in Ecology and Evolution, 2020, 8, .	1.1	11
29	Ecological traits and the spatial structure of competitive coexistence among carnivores. Ecology, 2020, 101, e03059.	1.5	61
30	Genomic approaches to identify hybrids and estimate admixture times in European wildcat populations. Scientific Reports, 2019, 9, 11612.	1.6	34
31	A metaproteomics approach reveals changes in mandibular lymph node microbiota of wild boar naturally exposed to an increasing trend of Mycobacterium tuberculosis complex infection. Tuberculosis, 2019, 114, 103-112.	0.8	2
32	Gastrointestinal parasite infestation in the alpine mountain hare (Lepus timidus varronis): Are abiotic environmental factors such as elevation, temperature and precipitation affecting prevalence of parasite species?. International Journal for Parasitology: Parasites and Wildlife, 2019, 9, 202-208.	0.6	8
33	The evolutionary history of the Cape hare (Lepus capensis sensu lato): insights for systematics and biogeography. Heredity, 2019, 123, 634-646.	1.2	12
34	Have the cake and eat it: Optimizing nondestructive DNA metabarcoding of macroinvertebrate samples for freshwater biomonitoring. Molecular Ecology Resources, 2019, 19, 863-876.	2.2	44
35	Feeding ecological knowledge: the underutilised power of faecal <scp>DNA</scp> approaches for carnivore diet analysis. Mammal Review, 2019, 49, 97-112.	2.2	60
36	Intraspecific genetic diversity and distribution of North African hedgehogs (Mammalia: Erinaceidae). Biological Journal of the Linnean Society, 2019, 127, 156-163.	0.7	5

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37	Glacial cycles drive rapid divergence of cryptic field vole species. Ecology and Evolution, 2019, 9, 14101-14113.	0.8	4
38	Mammals of Korea. Journal of Mammalogy, 2019, , .	0.6	0
39	Drivers of survival in a small mammal of conservation concern: An assessment using extensive genetic non-invasive sampling in fragmented farmland. Biological Conservation, 2019, 230, 131-140.	1.9	8
40	Does shortâ€term habitat management for the European rabbit (<scp>Oryctolagus) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 622 Td (cu	0.7	3
41	Red deer in Iberia: Molecular ecological studies in a southern refugium and inferences on European postglacial colonization history. PLoS ONE, 2019, 14, e0210282.	1.1	29
42	Insights into the evolution of the new variant rabbit haemorrhagic disease virus (GI.2) and the identification of novel recombinant strains. Transboundary and Emerging Diseases, 2018, 65, 983-992.	1.3	52
43	Winter color polymorphisms identify global hot spots for evolutionary rescue from climate change. Science, 2018, 359, 1033-1036.	6.0	91
44	Genome-wide associations identify novel candidate loci associated with genetic susceptibility to tuberculosis in wild boar. Scientific Reports, 2018, 8, 1980.	1.6	15
45	A genomic map of clinal variation across the European rabbit hybrid zone. Molecular Ecology, 2018, 27, 1457-1478.	2.0	30
46	Positive selection on the mitochondrialATP synthase 6and theNADH dehydrogenase 2genes across 22 hare species (genusLepus). Journal of Zoological Systematics and Evolutionary Research, 2018, 56, 428-443.	0.6	8
47	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
48	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
49	Epidemiology of RHDV2 (Lagovirus europaeus/GI.2) in free-living wild European rabbits in Portugal. Transboundary and Emerging Diseases, 2018, 65, e373-e382.	1.3	41
50	First genome-wide CNV mapping in FELIS CATUS using next generation sequencing data. BMC Genomics, 2018, 19, 895.	1.2	16
51	Combining molecular and landscape tools for targeting evolutionary processes in reserve design: An approach for islands. PLoS ONE, 2018, 13, e0200830.	1.1	7
52	Females know better: Sexâ€biased habitat selection by the European wildcat. Ecology and Evolution, 2018, 8, 9464-9477.	0.8	29
53	The effects of a motorway on movement behaviour and gene flow in a forest carnivore: Joint evidence from road mortality, radio tracking and genetics. Landscape and Urban Planning, 2018, 178, 217-227.	3.4	20
54	Combining genetic non-invasive sampling with spatially explicit capture-recapture models for density estimation of a patchily distributed small mammal. European Journal of Wildlife Research, 2018, 64, 1.	0.7	14

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55	Genetic non-invasive sampling (gNIS) as a cost-effective tool for monitoring elusive small mammals. European Journal of Wildlife Research, 2018, 64, 1.	0.7	45
56	Integrative approaches to guide conservation decisions: UsingÂgenomics to define conservation units and functionalÂcorridors. Molecular Ecology, 2018, 27, 3452-3465.	2.0	63
57	Recent range contractions in the globally threatened Pyrenean desman highlight the importance of stream headwater refugia. Animal Conservation, 2018, 21, 515-525.	1.5	15
58	Adaptive introgression underlies polymorphic seasonal camouflage in snowshoe hares. Science, 2018, 360, 1355-1358.	6.0	234
59	Range expansion underlies historical introgressive hybridization in the Iberian hare. Scientific Reports, 2017, 7, 40788.	1.6	35
60	Coccidiosis in European rabbit (Oryctolagus cuniculus algirus) populations in the Iberian Peninsula. Acta Parasitologica, 2017, 62, 229.	0.4	0
61	Wild opportunities with dedomestication genetics of rabbits. Restoration Ecology, 2017, 25, 330-332.	1.4	2
62	An update on the rabbit hemorrhagic disease virus (RHDV) strains circulating in Portugal in the 1990s: earliest detection of G3-G5 and G6. Archives of Virology, 2017, 162, 2061-2065.	0.9	3
63	Early-Onset Progressive Retinal Atrophy Associated with an IQCB1 Variant in African Black-Footed Cats (Felis nigripes). Scientific Reports, 2017, 7, 43918.	1.6	22
64	The transcriptional landscape of seasonal coat colour moult in the snowshoe hare. Molecular Ecology, 2017, 26, 4173-4185.	2.0	27
65	A FAS-ligand variant associated with autoimmune lymphoproliferative syndrome in cats. Mammalian Genome, 2017, 28, 47-55.	1.0	17
66	Precision Medicine in Cats: Novel Niemannâ€Pick Type C1 Diagnosed by Wholeâ€Genome Sequencing. Journal of Veterinary Internal Medicine, 2017, 31, 539-544.	0.6	30
67	Endemic species may have complex histories: withinâ€refugium phylogeography of an endangered Iberian vole. Molecular Ecology, 2017, 26, 951-967.	2.0	26
68	Characterization of old RHDV strains by complete genome sequencing identifies a novel genetic group. Scientific Reports, 2017, 7, 13599.	1.6	14
69	Mountain hare transcriptome and diagnostic markers as resources to monitor hybridization with European hares. Scientific Data, 2017, 4, 170178.	2.4	11
70	Urban Habitats Biodiversity Assessment (UrHBA): a standardized procedure for recording biodiversity and its spatial distribution in urban environments. Landscape Ecology, 2017, 32, 1753-1770.	1.9	15
71	Optimizing cameraâ€trapping protocols for characterizing mesocarnivore communities in southâ€western Europe. Journal of Zoology, 2017, 301, 23-31.	0.8	18
72	Ecotypes and evolutionary significant units in endangered North African gazelles. Biological Journal of the Linnean Society, 2017, 122, 286-300.	0.7	9

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73	Proposal for a unified classification system and nomenclature of lagoviruses. Journal of General Virology, 2017, 98, 1658-1666.	1.3	148
74	Spatial climate dynamics in the Iberian Peninsula since 15â€¯000â€¯yrâ€¯BP. Climate of the Past, 2016, 12, 1137-1149.	1.9	18
75	European wildcat populations are subdivided into five main biogeographic groups: consequences of Pleistocene climate changes or recent anthropogenic fragmentation?. Ecology and Evolution, 2016, 6, 3-22.	0.8	49
76	Disease-mediated bottom-up regulation: An emergent virus affects a keystone prey, and alters the dynamics of trophic webs. Scientific Reports, 2016, 6, 36072.	1.6	58
77	Coccidiosis in European rabbit (Oryctolagus cuniculus algirus) populations in the Iberian Peninsula. Acta Parasitologica, 2016, 61, 655.	0.4	0
78	Tuberculosis, genetic diversity and fitness in the red deer, Cervus elaphus. Infection, Genetics and Evolution, 2016, 43, 203-212.	1.0	19
79	Molecular and morphological insights into the origin of the invasive greater white-toothed shrew (Crocidura russula) in Ireland. Biological Invasions, 2016, 18, 857-871.	1.2	13
80	Genetic distinctiveness of the damselfly Coenagrion puella in North Africa: an overlooked and endangered taxon. Conservation Genetics, 2016, 17, 985-991.	0.8	12
81	LaGomiCsâ€”Lagomorph Genomics Consortium: An International Collaborative Effort for Sequencing the Genomes of an Entire Mammalian Order. Journal of Heredity, 2016, 107, 295-308.	1.0	19
82	Niche partitioning at the edge of the range: a multidimensional analysis with sympatric martens. Journal of Mammalogy, 2016, 97, 928-939.	0.6	28
83	Reply to Garner et al.. Trends in Ecology and Evolution, 2016, 31, 83-84.	4.2	24
84	Comparative Proteomics Identifies Host Immune System Proteins Affected by Infection with Mycobacterium bovis. PLoS Neglected Tropical Diseases, 2016, 10, e0004541.	1.3	12
85	Candidate genes underlying heritable differences in reproductive seasonality between wild and domestic rabbits. Animal Genetics, 2015, 46, 418-425.	0.6	14
86	Effect of microsatellite selection on individual and population genetic inferences: an empirical study using crossâ€specific and speciesâ€specific amplifications. Molecular Ecology Resources, 2015, 15, 747-760.	2.2	61
87	Biometrical analysis reveals major differences between the two subspecies of the European rabbit. Biological Journal of the Linnean Society, 2015, 116, 106-116.	0.7	18
88	Range dynamics driven by Quaternary climate oscillations explain the distribution of introgressed mt<scp>DNA</scp> ofÂLepus timidus origin in hares from the Iberian Peninsula. Journal of Biogeography, 2015, 42, 1727-1735.	1.4	21
89	Low persistence in nature of captive reared rabbits after restocking operations. European Journal of Wildlife Research, 2015, 61, 591-599.	0.7	9
90	Coccidiosis in European rabbit (Oryctolagus cuniculus algirus) populations in the Iberian Peninsula. Acta Parasitologica, 2015, 60, 350-5.	0.4	7

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91	Can we predict habitat quality from space? A multi-indicator assessment based on an automated knowledge-driven system. International Journal of Applied Earth Observation and Geoinformation, 2015, 37, 106-113.	1.4	19
92	Is the New Variant RHDV Replacing Genogroup 1 in Portuguese Wild Rabbit Populations?. Viruses, 2015, 7, 27-36.	1.5	66
93	Toward a genome-wide approach for detecting hybrids: informative SNPs to detect introgression between domestic cats and European wildcats (Felis silvestris). Heredity, 2015, 115, 195-205.	1.2	51
94	Local extinctions and range contraction of the endangeredCoenagrion mercurialein North Africa. International Journal of Odonatology, 2015, 18, 137-152.	0.5	9
95	Genomics and the challenging translation into conservation practice. Trends in Ecology and Evolution, 2015, 30, 78-87.	4.2	469
96	Feline mitochondrial DNA sampling for forensic analysis: When enough is enough!. Forensic Science International: Genetics, 2015, 16, 52-57.	1.6	9
97	Combining distribution modelling and non-invasive genetics to improve range shift forecasting. Ecological Modelling, 2015, 297, 171-179.	1.2	16
98	Genetic identification of endangered <scp>N</scp>orth <scp>A</scp>frican ungulates using noninvasive sampling. Molecular Ecology Resources, 2015, 15, 652-661.	2.2	25
99	Ecological interactions and species coexistence in Iberian mesocarnivore communities - Extended summary and main results.. Galemys Spanish Journal of Mammalogy, 2015, 27, 47-57.	0.2	7
100	The impact of management practices and past demographic history on the genetic diversity of red deer (Cervus elaphus): an assessment of population and individual fitness. Biological Journal of the Linnean Society, 2014, 111, 209-223.	0.7	23
101	Detection of RHDV strains in the Iberian hare (Lepus granatensis): earliest evidence of rabbit lagovirus cross-species infection. Veterinary Research, 2014, 45, 94.	1.1	24
102	The Elusive Nature of Adaptive Mitochondrial DNA Evolution of an Arctic Lineage Prone to Frequent Introgression. Genome Biology and Evolution, 2014, 6, 886-896.	1.1	78
103	Experimental study on the effect of cover and vaccination on the survival of juvenile European rabbits. Population Ecology, 2014, 56, 195-202.	0.7	3
104	Efficiency of hair snares and camera traps to survey mesocarnivore populations. European Journal of Wildlife Research, 2014, 60, 279-289.	0.7	29
105	Evidence for niche similarities in the allopatric sister species <scp>L</scp>epus castroviejoi and <scp>L</scp>epus corsicanus. Journal of Biogeography, 2014, 41, 977-986.	1.4	17
106	Mitochondrial phylogeography of the European wild boar: the effect of climate on genetic diversity and spatial lineage sorting across Europe. Journal of Biogeography, 2014, 41, 987-998.	1.4	56
107	Conservation implications of the evolutionary history and genetic diversity hotspots of the snowshoe hare. Molecular Ecology, 2014, 23, 2929-2942.	2.0	32
108	Home-loving boreal hare mitochondria survived several invasions in Iberia: the relative roles of recurrent hybridisation and allele surfing. Heredity, 2014, 112, 265-273.	1.2	30

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109	Molecular phylogeny of the Western PalaearcticCordulegastertaxa (Odonata: Anisoptera:) Tj ETQq1 1 0.784314 rgBT /Overloc	0.7	19
110	The hidden history of the snowshoe hare, <scp>L</scp>epus americanus: extensive mitochondrial <scp>DNA</scp> introgression inferred from multilocus genetic variation. Molecular Ecology, 2014, 23, 4617-4630.	2.0	40
111	Sequencing of Sylvilagus VDJ genes reveals a new VHa allelic lineage and shows that ancient VH lineages were retained differently in leporids. Immunogenetics, 2014, 66, 719-726.	1.2	6
112	Molecular and ecological signs of mitochondrial adaptation: consequences for introgression?. Heredity, 2014, 113, 277-286.	1.2	37
113	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
114	Colonization history of Mallorca Island by the European rabbit,Oryctolagus cuniculus, and the Iberian hare,Lepus granatensis(Lagomorpha: Leporidae). Biological Journal of the Linnean Society, 2014, 111, 748-760.	0.7	7
115	Leporid immunoglobulin G shows evidence of strong selective pressure on the hinge and CH3 domains. Open Biology, 2014, 4, 140088.	1.5	18
116	A Critically Endangered new dragonfly species from Morocco: Onychogomphus boudoti sp. nov. (Odonata: Gomphidae). Zootaxa, 2014, 3856, 349-65.	0.2	19
117	Detection of RHDV strains in the Iberian hare (Lepus granatensis): earliest evidence of rabbit lagovirus cross-species infection. Veterinary Research, 2014, 45, 94.	1.1	24
118	Land Bridge Calibration of Rates of Molecular Evolution in a Widespread Rodent. , 2014, , 69-86.		3
119	Genetic structure of wildcat (Felis silvestris) populations in Italy. Ecology and Evolution, 2013, 3, 2443-2458.	0.8	58
120	Adapted conservation measures are required to save the Iberian lynx in a changing climate. Nature Climate Change, 2013, 3, 899-903.	8.1	96
121	Estimating homeâ€range size: when to include a third dimension?. Ecology and Evolution, 2013, 3, 2285-2295.	0.8	12
122	Catch Me If You Can: Diel Activity Patterns of Mammalian Prey and Predators. Ethology, 2013, 119, 1044-1056.	0.5	128
123	Genetic identification of <scp>I</scp>berian rodent species using both mitochondrial and nuclear loci: application to noninvasive sampling. Molecular Ecology Resources, 2013, 13, 43-56.	2.2	55
124	Factors affecting the (in)accuracy of mammalian mesocarnivore scat identification in <scp>S</scp>outhâ€western <scp>E</scp>urope. Journal of Zoology, 2013, 289, 243-250.	0.8	48
125	Sequencing of modern Lepus VDJ genes shows that the usage of VHn genes has been retained in both Oryctolagus and Lepus that diverged 12 million years ago. Immunogenetics, 2013, 65, 777-784.	1.2	18
126	Reference-Free Population Genomics from Next-Generation Transcriptome Data and the Vertebrateâ€“Invertebrate Gap. PLoS Genetics, 2013, 9, e1003457.	1.5	157

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127	New Variant of Rabbit Hemorrhagic Disease Virus, Portugal, 2012â€“2013. Emerging Infectious Diseases, 2013, 19, 1900-2.	2.0	86
128	Cryptic speciation in the field vole: a multilocus approach confirms three highly divergent lineages in <scp>E</scp>urasia. Molecular Ecology, 2012, 21, 6015-6032.	2.0	59
129	Recurrent Introgression of Mitochondrial DNA Among Hares (Lepus spp.) Revealed by Species-Tree Inference and Coalescent Simulations. Systematic Biology, 2012, 61, 367.	2.7	111
130	Giant sex chromosomes retained within the Portuguese lineage of the field vole (Microtus agrestis). Acta Theriologica, 2012, 57, 377-382.	1.1	6
131	Past, Present and Future Distributions of an Iberian Endemic, Lepus granatensis: Ecological and Evolutionary Clues from Species Distribution Models. PLoS ONE, 2012, 7, e51529.	1.1	31
132	Parapatric species and the implications for climate change studies: a case study on hares in <scp>E</scp>urope. Global Change Biology, 2012, 18, 1509-1519.	4.2	49
133	Genetic Diversity of MaghrebianHottentotta(Scorpiones: Buthidae) Scorpions Based on CO1: New Insights on the Genus Phylogeny and Distribution. African Invertebrates, 2011, 52, 135-143.	0.5	27
134	Evaluation of attractants for non-invasive studies of Iberian carnivore communities. Wildlife Research, 2011, 38, 446.	0.7	45
135	INTERSPECIFIC X-CHROMOSOME AND MITOCHONDRIAL DNA INTROGRESSION IN THE IBERIAN HARE: SELECTION OR ALLELE SURFING?. Evolution; International Journal of Organic Evolution, 2011, 65, 1956-1968.	1.1	29
136	Hares in Corsica: high prevalence of Lepus corsicanus and hybridization with introduced L. europaeus and L. granatensis. European Journal of Wildlife Research, 2011, 57, 313-321.	0.7	19
137	Molecular bases of genetic diversity and evolution of the immunoglobulin heavy chain variable region (IGHV) gene locus in leporids. Immunogenetics, 2011, 63, 397-408.	1.2	31
138	Introgression of mitochondrial DNA among Myodes voles: consequences for energetics?. BMC Evolutionary Biology, 2011, 11, 355.	3.2	50
139	Species identification using a small nuclear gene fragment: application to sympatric wild carnivores from South-western Europe. Conservation Genetics, 2010, 11, 1023-1032.	0.8	36
140	Genetic diversity of wild boar populations and domestic pig breeds (Sus scrofa) in South-western Europe. Biological Journal of the Linnean Society, 2010, 101, 797-822.	0.7	42
141	The usefulness of field data and hunting statistics in the assessment of wild rabbit (Oryctolagus) Tj ETQq1 1 0.784314 rgBT /Overlock	0.7	21
142	Genetic diversity within scorpions of the genus Buthus from the Iberian Peninsula: mitochondrial DNA sequence data indicate additional distinct cryptic lineages. Journal of Arachnology, 2010, 38, 206-211.	0.3	28
143	Evolution of rabbit haemorrhagic disease virus (RHDV) in the European rabbit (Oryctolagus) Tj ETQq1 1 0.784314 rgBT /Overlock 10	0.8	50
144	Influence of habitat management on the abundance and diet of wild rabbit (Oryctolagus cuniculus) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5 487-496.	0.7	55

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145	Phylogeography of the brown hare (Lepus europaeus) in Europe: a legacy of southâ€eastern Mediterranean refugia?. Journal of Biogeography, 2009, 36, 515-528.	1.4	63
146	The genomic legacy from the extinct Lepus timidus to the three hare species of Iberia: contrast between mtDNA, sex chromosomes and autosomes. Molecular Ecology, 2009, 18, 2643-2658.	2.0	69
147	Spatial ecology of the European wildcat in a Mediterranean ecosystem: dealing with small radioâ€tracking datasets in species conservation. Journal of Zoology, 2009, 279, 27-35.	0.8	89
148	Field experimental vaccination campaigns against myxomatosis and their effectiveness in the wild. Vaccine, 2009, 27, 6998-7002.	1.7	24
149	Advancing ecological understandings through technological transformations in noninvasive genetics. Molecular Ecology Resources, 2009, 9, 1279-1301.	2.2	296
150	Molecular analysis of hybridisation between wild and domestic cats (Felis silvestris) in Portugal: implications for conservation. Conservation Genetics, 2008, 9, 1-11.	0.8	100
151	Diet of the Iberian hare (Lepus granatensis) in a mountain ecosystem. European Journal of Wildlife Research, 2008, 54, 571-579.	0.7	21
152	Evidence of autumn reproduction in female European hares (Lepus europaeus) from southern Europe. European Journal of Wildlife Research, 2008, 54, 581-587.	0.7	12
153	Evidence for genetic similarity of two allopatric European hares (Lepus corsicanus and L.) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 5 1191-1197.	1.2	39
154	Genetic diversity within Scorpio maurus (Scorpiones: Scorpionidae) from morocco: Preliminary evidence based on CO1 mitochondrial DNA sequences. Biologia (Poland), 2008, 63, 1157-1160.	0.8	18
155	Population genetics of cape and brown hares (Lepus capensis and L. europaeus): A test of Petter's hypothesis of conspecificity. Biochemical Systematics and Ecology, 2008, 36, 22-39.	0.6	43
156	Hybridization versus conservation: are domestic cats threatening the genetic integrity of wildcats (Felis silvestris silvestris) in Iberian Peninsula?. Philosophical Transactions of the Royal Society B: Biological Sciences, 2008, 363, 2953-2961.	1.8	91
157	Endemic Sand Dune Vegetation of the Northwest Iberian Peninsula: Diversity, Dynamics, and Significance for Bioindication and Monitoring of Coastal Landscapes. Journal of Coastal Research, 2008, 2, 113-121.	0.1	29
158	The ubiquitous mountain hare mitochondria: multiple introgressive hybridization in hares, genusLepus. Philosophical Transactions of the Royal Society B: Biological Sciences, 2008, 363, 2831-2839.	1.8	111
159	Overview of Lagomorph Research: What we have learned and what we still need to do. , 2008, , 381-391.		8
160	Patterns of genetic diversity within and between Myotis d. daubentonii and M. d. nathalinae derived from cytochromebmtDNA sequence data. Acta Chiropterologica, 2007, 9, 379-389.	0.2	7
161	Evolutionary relationships among hares from North Africa (Lepus sp. or Lepus spp.), cape hares (L.) Tj ETQq1 1 0.784314 rgBT /Overlo allozyme data. Journal of Zoological Systematics and Evolutionary Research, 2006, 44, 88-99.	0.6	21
162	The rise and fall of the mountain hare (Lepus timidus) during Pleistocene glaciations: expansion and retreat with hybridization in the Iberian Peninsula. Molecular Ecology, 2006, 16, 605-618.	2.0	95

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163	Hares on thin ice: Introgression of mitochondrial DNA in hares and its implications for recent phylogenetic analyses. Molecular Phylogenetics and Evolution, 2006, 40, 640-641.	1.2	40
164	Invasion from the cold past: extensive introgression of mountain hare (Lepus timidus) mitochondrial DNA into three other hare species in northern Iberia. Molecular Ecology, 2005, 14, 2459-2464.	2.0	183
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170	Hotspot variation at the CH2-CH3 interface of leporid IgG antibodies (Oryctolagus, Sylvilagus and) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50	1.2	10
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