

# JosÃ© Antonio Blanco-Aguiar

## List of Publications by Year in descending order

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Version: 2024-02-01

33  
papers

999  
citations

516215

16  
h-index

454577

30  
g-index

34  
all docs

34  
docs citations

34  
times ranked

1575  
citing authors

#	ARTICLE	IF	CITATIONS
1	Can we model distribution of population abundance from wildlifeâ€œvehicles collision data?. Ecography, 2022, 2022, .	2.1	12
2	Revisiting wild boar spatial models based on hunting yields to assess their predictive performance on interpolation and extrapolation areas. Ecological Modelling, 2022, 471, 110041.	1.2	2
3	Analysis of wild boarâ€œdomestic pig interface in Europe: spatial overlapping and fine resolution approach in several countries. EFSA Supporting Publications, 2021, 18, 1995E.	0.3	7
4	A loss-of-function mutation in RORB disrupts saltatorial locomotion in rabbits. PLoS Genetics, 2021, 17, e1009429.	1.5	10
5	Update of model for wild boar abundance based on hunting yield and first models based on occurrence for wild ruminants at European scale. EFSA Supporting Publications, 2021, 18, 6825E.	0.3	5
6	Brain Transcriptomics of Wild and Domestic Rabbits Suggests That Changes in Dopamine Signaling and Ciliary Function Contributed to Evolution of Tameness. Genome Biology and Evolution, 2020, 12, 1918-1928.	1.1	17
7	Improving models of wild boar hunting yield distribution: new insights for predictions at fine spatial resolution. EFSA Supporting Publications, 2020, 17, 1980E.	0.3	1
8	Update of occurrence and hunting yieldâ€œbased data models for wild boar at European scale: new approach to handle the bioregion effect. EFSA Supporting Publications, 2020, 17, 1871E.	0.3	6
9	Analysis of wild boarâ€œdomestic pig interface in Europe: preliminary analysis. EFSA Supporting Publications, 2020, 17, 1834E.	0.3	6
10	Applying the Darwin core standard to the monitoring of wildlife species, their management and estimated records. EFSA Supporting Publications, 2020, 17, 1841E.	0.3	9
11	Harmonization of the use of hunting statistics for wild boar density estimation in different study areas. EFSA Supporting Publications, 2019, 16, 1706E.	0.3	14
12	Science-based wildlife disease response. Science, 2019, 364, 943-944.	6.0	42
13	A genomic map of clinal variation across the European rabbit hybrid zone. Molecular Ecology, 2018, 27, 1457-1478.	2.0	30
14	Analysis of hunting statistics collection frameworks for wild boar across Europe and proposals for improving the harmonisation of data collection. EFSA Supporting Publications, 2018, 15, 1523E.	0.3	10
15	First assessment of the potential introduction by hunters of eastern cottontail rabbits ( <i>Sylvilagus</i> ) Tj ETQq1 1 0.784314 rgBT <sub>5</sub> /Overlo	0.7	5
16	Changes in brain architecture are consistent with altered fear processing in domestic rabbits. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 7380-7385.	3.3	45
17	Full genome sequences are key to disclose RHDV2 emergence in the Macaronesian islands. Virus Genes, 2018, 54, 1-4.	0.7	9
18	Dwarfism and Altered Craniofacial Development in Rabbits Is Caused by a 12.1 kb Deletion at the <i>HMGA2</i> Locus. Genetics, 2017, 205, 955-965.	1.2	30

#	ARTICLE	IF	CITATIONS
19	Climatic and geographic effects on the spatial genetic pattern of a landbird species ( <i>Alectoris</i> ) Tj ETQq1 1 0.784314 rgBT /Overloc	0.7	18
20	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
21	The Genomic Architecture of Population Divergence between Subspecies of the European Rabbit. PLoS Genetics, 2014, 10, e1003519.	1.5	82
22	A Comparison of Brain Gene Expression Levels in Domesticated and Wild Animals. PLoS Genetics, 2012, 8, e1002962.	1.5	130
23	Evidence for Widespread Positive and Purifying Selection Across the European Rabbit ( <i>Oryctolagus</i> ) Tj ETQq1 1 0.784314 rgBT /Overloc	3.5	71
24	Is the interaction between rabbit hemorrhagic disease and hyperpredation by raptors a major cause of the red-legged partridge decline in Spain?. European Journal of Wildlife Research, 2012, 58, 433-439.	0.7	20
25	Phylogeography and genetic structure of the red-legged partridge ( <i>Alectoris rufa</i> ): more evidence for refugia within the Iberian glacial refugium. Molecular Ecology, 2011, 20, 2628-2642.	2.0	30
26	SPECIATION IN THE EUROPEAN RABBIT ( <i>ORYCTOLAGUS CUNICULUS</i> ): ISLANDS OF DIFFERENTIATION ON THE X CHROMOSOME AND AUTOSOMES. Evolution; International Journal of Organic Evolution, 2010, 64, 3443-3460.	1.1	71
27	Survival and causes of mortality among wild Red-legged Partridges ( <i>Alectoris rufa</i> ) in southern Spain: implications for conservation. Ibis, 2009, 151, 720-730.	1.0	42
28	Habitat selection and home range size of red-legged partridges in Spain. Agriculture, Ecosystems and Environment, 2008, 126, 158-162.	2.5	50
29	Assessment of game restocking contributions to anthropogenic hybridization: the case of the Iberian red-legged partridge. Animal Conservation, 2008, 11, 535-545.	1.5	92
30	Sixteen new polymorphic microsatellite markers isolated for red-legged partridge ( <i>Alectoris rufa</i> ) and related species. Molecular Ecology Notes, 2007, 7, 1349-1351.	1.7	11
31	Evolution of life history traits in Leporidae: a test of nest predation and seasonality hypotheses. Biological Journal of the Linnean Society, 2006, 88, 603-610.	0.7	13
32	Food habits of European badgers ( <i>Meles meles</i> ) along an altitudinal gradient of Mediterranean environments: a field test of the earthworm specialization hypothesis. Canadian Journal of Zoology, 2004, 82, 41-51.	0.4	50
33	Spatial variation in helminth community structure in the red-legged partridge ( <i>Alectoris rufa</i> ): effects of definitive host density. Parasitology, 2004, 129, 101-113.	0.7	56