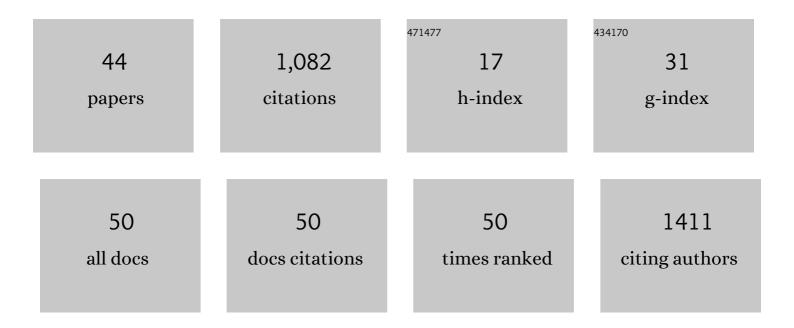
Laura Iacolina

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/241330/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	New developments in the field of genomic technologies and their relevance to conservation management. Conservation Genetics, 2022, 23, 217-242.	1.5	26
2	Resilience to Historical Human Manipulations in the Genomic Variation of Italian Wild Boar Populations. Frontiers in Ecology and Evolution, 2022, 10, .	2.2	6
3	Main roads and land cover shaped the genetic structure of a Mediterranean island wild boar population. Ecology and Evolution, 2022, 12, e8804.	1.9	4
4	Past, present and future of chamois science. Wildlife Biology, 2022, 2022, .	1.4	6
5	Comparing DNA metabarcoding with faecal analysis for diet determination of the Eurasian otter (Lutra lutra) in Vejlerne, Denmark. Mammal Research, 2021, 66, 115-122.	1.3	13
6	eDNA metabarcoding for biodiversity assessment, generalist predators as sampling assistants. Scientific Reports, 2021, 11, 6820.	3.3	20
7	12. Conclusions. , 2021, , 305-310.		0
8	1. African swine fever (ASF), the pig health challenge of the century. , 2021, , 11-24.		2
9	A Mother's Story, Mitogenome Relationships in the Genus Rupicapra. Animals, 2021, 11, 1065.	2.3	6
10	Genome-wide profiles indicate wolf population connectivity within the eastern Carpathian Mountains. Genetica, 2020, 148, 33-39.	1.1	3
11	A refined genome-wide association study of posthitis in lowland BiaÅ,owieza population of the European bison (Bison bonasus). European Journal of Wildlife Research, 2020, 66, 1.	1.4	7
12	MHC Genotyping by SSCP and Amplicon-Based NGS Approach in Chamois. Animals, 2020, 10, 1694.	2.3	2
13	Fragmentation and Translocation Distort the Genetic Landscape of Ungulates: Red Deer in the Netherlands. Frontiers in Ecology and Evolution, 2020, 8, .	2.2	15
14	eDNA and metabarcoding for rewilding projects monitoring, a dietary approach. Mammalian Biology, 2020, 100, 411-418.	1.5	6
15	Microsatellite based assignment reveals history of extirpated mountain ungulate. Genetica, 2020, 148, 41-46.	1.1	1
16	Conservation Genomic Analysis of the Croatian Indigenous Black Slavonian and Turopolje Pig Breeds. Frontiers in Genetics, 2020, 11, 261.	2.3	17
17	Genomic variability in the extinct steppe bison (Bison priscus) compared to the European bison (Bison) Tj ETQq1	l 0.78431 1.3	4 ₄ gBT /Ove
18	Genomic analyses suggest adaptive differentiation of northern European native cattle breeds.	3.1	12

Evolutionary Applications, 2019, 12, 1096-1113.

LAURA IACOLINA

#	Article	IF	CITATIONS
19	Rewilding and conservation genomics: How developments in (re)colonization ecology and genomics can offer mutual benefits for understanding contemporary evolution. Global Ecology and Conservation, 2019, 17, e00502.	2.1	7
20	Hybridisation in European ungulates: an overview of the current status, causes, and consequences. Mammal Review, 2019, 49, 45-59.	4.8	60
21	The use of museum skins for genomic analyses of temporal genetic diversity in wild species. Conservation Genetics Resources, 2019, 11, 499-503.	0.8	3
22	Hotspots of recent hybridization between pigs and wild boars in Europe. Scientific Reports, 2018, 8, 17372.	3.3	53
23	How much does it cost to look like a pig in a wild boar group?. Behavioural Processes, 2017, 138, 123-126.	1.1	15
24	Genome-wide SNP data unveils the globalization of domesticated pigs. Genetics Selection Evolution, 2017, 49, 71.	3.0	114
25	Novel Graphical Analyses of Runs of Homozygosity among Species and Livestock Breeds. International Journal of Genomics, 2016, 2016, 1-8.	1.6	26
26	Lack of polymorphism at the MC1R wild-type allele and evidence of domestic allele introgression across European wild boar populations. Mammalian Biology, 2016, 81, 477-479.	1.5	17
27	Novel Yâ€chromosome short tandem repeats in <i>Sus scrofa</i> and their variation in European wild boar and domestic pig populations. Animal Genetics, 2016, 47, 682-690.	1.7	6
28	Genomic diversity and differentiation of a managed island wild boar population. Heredity, 2016, 116, 60-67.	2.6	41
29	Prospects and challenges for the conservation of farm animal genomic resources, 2015-2025. Frontiers in Genetics, 2015, 6, 314.	2.3	64
30	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.	3.0	56
31	Are captive wild boar more introgressed than free-ranging wild boar? Two case studies in Italy. European Journal of Wildlife Research, 2014, 60, 459-467.	1.4	21
32	Influence of management regime and population history on genetic diversity and population structure of brown hares (Lepus europaeus) in an Italian province. European Journal of Wildlife Research, 2013, 59, 783-793.	1.4	9
33	Differentiation under isolation and genetic structure of Sardinian hares as revealed by craniometric analysis, mitochondrial DNA and microsatellites. Journal of Zoological Systematics and Evolutionary Research, 2012, 50, 328-337.	1.4	9
34	Genetic diversity in the European wild boar Sus scrofa: phylogeography, population structure and wild x domestic hybridization. Mammal Review, 2011, 41, 125-137.	4.8	106
35	Fine-scale genetic structure suggests low levels of short-range gene flow in a wolf population of the Italian Apennines. European Journal of Wildlife Research, 2011, 57, 949-958.	1.4	23
36	Effects of human perturbation on the genetic make-up of an island population: the case of the Sardinian wild boar. Heredity, 2011, 106, 1012-1020.	2.6	45

LAURA IACOLINA

#	Article	IF	CITATIONS
37	Current status of the Sardinian partridge (Alectoris barbara) assessed by molecular markers. European Journal of Wildlife Research, 2010, 56, 33-42.	1.4	17
38	Y-chromosome microsatellite variation in Italian wolves: A contribution to the study of wolf-dog hybridization patterns. Mammalian Biology, 2010, 75, 341-347.	1.5	38
39	Nonkin Associations in Wild Boar Social Units. Journal of Mammalogy, 2009, 90, 666-674.	1.3	25
40	Ancient vs. recent processes as factors shaping the genetic variation of the European wild boar: are the effects of the last glaciation still detectable?. Molecular Ecology, 2008, 17, 1745-1762.	3.9	129
41	Mitochondrial CR-1 Variation in Sardinian Hares and Its Relationships with Other Old World Hares (Genus Lepus). Biochemical Genetics, 2007, 45, 305-323.	1.7	14
42	An empirical approach for reliable microsatellite genotyping of wolf DNA from multiple noninvasive sources. Conservation Genetics, 2006, 7, 813-823.	1.5	29
43	The Balkan chamois, an archipelago or a peninsula? Insights from nuclear and mitochondrial DNA. Conservation Genetics, 0, , 1.	1.5	2
44	Association between dietary nutrient intake and sarcopenia in older adults. , 0, , 215-229.		0