Analyses of Possible Domestic Pig Contribution in Two Boar

Acta Agriculturae Scandinavica - Section A: Animal Science 53, 161-165

DOI: 10.1080/09064710310010602

Citation Report

#	Article	IF	CITATIONS
1	Nuclear and mitochondrial evolutionary analyses of Collared, White-lipped, and Chacoan peccaries (Tayassuidae). Molecular Phylogenetics and Evolution, 2005, 34, 181-189.	1.2	30
2	Mitochondrial DNA sequence variations in some Italian wild boar populations. Journal of Animal Breeding and Genetics, 2009, 126, 154-163.	0.8	7
3	Mitochondrial lineages reveal intense gene flow between Iberian wild boars and South Iberian pig breeds. Animal Genetics, 2012, 43, 35-41.	0.6	24
4	The Balkans and the colonization of Europe: the postâ€glacial range expansion of the wild boar, <i>Sus scrofa</i> . Journal of Biogeography, 2012, 39, 713-723.	1.4	64
5	Genetic evidence for introgression between domestic pigs and wild boars (<i>Sus scrofa</i>) in Belgium and Luxembourg: a comparative approach with multiple marker systems. Biological Journal of the Linnean Society, 2013, 110, 104-115.	0.7	41
6	Wild boar meat sensory attributes contributing general meat quality. Bulletin of the Veterinary Institute in Pulawy = Biuletyn Instytutu Weterynarii W Pulawach, 2013, 57, 357-363.	0.4	6
7	Mitochondrial diversity supports multiple origins for invasive pigs. Journal of Wildlife Management, 2014, 78, 202-213.	0.7	19
8	Muscle fibre characteristics, enzyme activity and meat colour of wild boar (Sus scrofa s. L.) muscle with 2n=36 compared to those of phenotypically similar crossbreeds (2n=37 and 2n=38). Meat Science, 2014, 98, 272-278.	2.7	7
9	Size matters: A comparative analysis of pig domestication. Holocene, 2016, 26, 327-332.	0.9	13
10	Evolutionary Relationships and Taxonomy of Suidae and Tayassuidae. , 0, , 1-19.		4
11	High domestic pig contribution to the local gene pool of free-living European wild boar: a case study in Poland. Mammal Research, 2018, 63, 65-71.	0.6	14
13	Patterns of genetic variation on wild pig (<i>Sus scrofa</i>) populations over a complete range of the species in Argentina. Mammalia, 2022, 86, 359-372.	0.3	5