Maja FerenÄaković

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/10591085/publications.pdf

Version: 2024-02-01

840776 1058476 14 922 11 14 citations h-index g-index papers 14 14 14 1021 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Genomic Characterization of the Three Balkan Livestock Guardian Dogs. Sustainability, 2021, 13, 2289.	3.2	4
2	Conservation Genomic Analysis of the Croatian Indigenous Black Slavonian and Turopolje Pig Breeds. Frontiers in Genetics, 2020, $11,261$.	2.3	17
3	Hypocalcemia After Completion Thyroidectomy for Papillary Thyroid Carcinoma. Acta Clinica Croatica, 2020, 59, 136-145.	0.2	1
4	AUTALASSO: an automatic adaptive LASSO for genome-wide prediction. BMC Bioinformatics, 2019, 20, 167.	2.6	20
5	Misidentification of runs of homozygosity islands in cattle caused by interference with copy number variation or large intermarker distances. Genetics Selection Evolution, 2018, 50, 43.	3.0	32
6	Genomic characterization of Pinzgau cattle: genetic conservation and breeding perspectives. Conservation Genetics, 2017, 18, 893-910.	1.5	39
7	Genome-wide mapping and estimation of inbreeding depression of semen quality traits in a cattle population. Journal of Dairy Science, 2017, 100, 4721-4730.	3.4	89
8	Genomic dissection of inbreeding depression: a gate to new opportunities. Revista Brasileira De Zootecnia, 2017, 46, 773-782.	0.8	26
9	GENOMIC VARIABILITY AMONG CATTLE POPULATIONS BASED ON RUNS OF HOMOZYGOSITY. Poljoprivreda, 2015, 21, 44-47.	0.5	5
10	Genomic analysis for managing small and endangered populations: a case study in Tyrol Grey cattle. Frontiers in Genetics, 2015, 6, 173.	2.3	46
11	Prospects and challenges for the conservation of farm animal genomic resources, 2015-2025. Frontiers in Genetics, 2015, 6, 314.	2.3	64
12	Assessment of autozygosity in Nellore cows (Bos indicus) through high-density SNP genotypes. Frontiers in Genetics, 2015, 6, 5.	2.3	69
13	Inbreeding and runs of homozygosity: A possible solution to an old problem. Livestock Science, 2014, 166, 26-34.	1.6	283
14	Estimating autozygosity from high-throughput information: effects of SNP density and genotyping errors. Genetics Selection Evolution, 2013, 45, 42.	3.0	227