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List of Publications by Year in descending order

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Version: 2024-02-01

15
papers

434
citations

1478505

6
h-index

996975

15
g-index

15
all docs

15
docs citations

15
times ranked

595
citing authors

#	ARTICLE	IF	CITATIONS
1	Genomic scan of selective sweeps in thin and fat tail sheep breeds for identifying of candidate regions associated with fat deposition. <i>BMC Genetics</i> , 2012, 13, 10.	2.7	236
2	Predictive performance of genomic selection methods for carcass traits in Hanwoo beef cattle: impacts of the genetic architecture. <i>Genetics Selection Evolution</i> , 2017, 49, 1.	3.0	89
3	Comparison of conventional BLUP and single-step genomic BLUP evaluations for yearling weight and carcass traits in Hanwoo beef cattle using single trait and multi-trait models. <i>PLoS ONE</i> , 2019, 14, e0223352.	2.5	24
4	Nutritional and physiological responses of broiler chickens to dietary supplementation with de-oiled soyabean lecithin at different metabolisable energy levels and various fat sources. <i>British Journal of Nutrition</i> , 2019, 122, 863-872.	2.3	16
5	De-oiled soy lecithin positively influenced growth performance, nutrient digestibility, histological intestinal alteration, and antioxidant status in turkeys fed with low energy diets. <i>British Poultry Science</i> , 2021, 62, 858-867.	1.7	11
6	Weighted Single-Step Genome-Wide Association Study Uncovers Known and Novel Candidate Genomic Regions for Milk Production Traits and Somatic Cell Score in Valle del Belice Dairy Sheep. <i>Animals</i> , 2022, 12, 1155.	2.3	10
7	Synchronous feeding of liquid protein source with different grains on performance, digestibility, ruminal fermentation, blood metabolites, and carcass characters in growing lambs. <i>Tropical Animal Health and Production</i> , 2020, 52, 829-837.	1.4	8
8	Genome-wide identification of copy number variation and association with fat deposition in thin and fat-tailed sheep breeds. <i>Scientific Reports</i> , 2022, 12, .	3.3	7
9	Genome-Wide Association Study of Weaning Traits in Lori-Bakhtiari Sheep. <i>Annals of Animal Science</i> , 2020, 20, 811-824.	1.6	6
10	Hitchhiking Mapping of Candidate Regions Associated with Fat Deposition in Iranian Thin and Fat Tail Sheep Breeds Suggests New Insights into Molecular Aspects of Fat Tail Selection. <i>Animals</i> , 2022, 12, 1423.	2.3	6
11	Genome-wide association study for postweaning weight traits in Lori-Bakhtiari sheep. <i>Tropical Animal Health and Production</i> , 2021, 53, 163.	1.4	5
12	Genome-wide selection of discriminant SNP markers for breed assignment in indigenous sheep breeds. <i>Annals of Animal Science</i> , 2021, 21, 807-831.	1.6	5
13	Effects of a Grain Source (Corn Versus Barley) and Starter Protein Content on Performance, Ruminal Fermentation, and Blood Metabolites in Holstein Dairy Calves. <i>Animals</i> , 2020, 10, 1722.	2.3	4
14	Genome-Wide Association Study Using Fix-Length Haplotypes and Network Analysis Revealed New Candidate Genes for Nematode Resistance and Body Weight in Blackface Lambs. <i>Annals of Animal Science</i> , 2020, 20, 445-464.	1.6	4
15	Single and combined effects of phytase and citric acid on growth performance, nutrient digestibility, bone characteristics, intestinal morphology, and blood components in meat-type quails fed low-phosphorous diets. <i>Animal Feed Science and Technology</i> , 2020, 269, 114677.	2.2	3