Linsheng Gui

List of Publications by Year in descending order

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		1478505	1372567
10	157	6	10
papers	citations	h-index	g-index
10	10	10	157
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Associations between UASMS2 polymorphism in leptin gene and growth, carcass and meat quality traits of cattle: a meta-analysis. Animal Biotechnology, 2022, 33, 279-288.	1.5	7
2	Genetic variants in the serum amyloid A2 (<i>SAA2</i>) gene as a potential marker for milk production traits in Chinese Holstein cows. Veterinary Medicine and Science, 2022, 8, 1835-1840.	1.6	1
3	CEBPÎ 2 binding directly to the promoter region drives CEBPÉ' transcription and improves FABP4 transcriptional activity in adipose tissue of yak (Bos grunniens). Research in Veterinary Science, 2021, 141, 174-179.	1.9	4
4	Eukaryotic expression, Co-IP and MS identify BMPR-1B protein–protein interaction network. Biological Research, 2020, 53, 24.	3.4	15
5	Bioinformatics analysis and genetic polymorphisms in genomic region of the bovine <i>SH2B2</i> gene and their associations with molecular breeding for body size traits in qinchuan beef cattle. Bioscience Reports, 2020, 40, .	2.4	16
6	Association of polymorphisms in bone morphogenetic protein receptor-1B gene exon-9 with litter size in Dorset, Mongolian, and Small Tail Han ewes. Asian-Australasian Journal of Animal Sciences, 2019, 32, 949-955.	2.4	10
7	Composition, diversity and function of gastrointestinal microbiota in wild red-billed choughs (Pyrrhocorax pyrrhocorax). International Microbiology, 2019, 22, 491-500.	2.4	15
8	The effect of haplotypes in the promoter region of SIRT4 gene on the ultrasound traits in Qinchuan cattle. Tropical Animal Health and Production, 2019, 51, 1877-1882.	1.4	7
9	Association between FASN gene polymorphisms ultrasound carcass traits and intramuscular fat in Qinchuan cattle. Gene, 2018, 645, 55-59.	2.2	54
10	Haplotype distribution in the class I sirtuin genes and their associations with ultrasound carcass traits in Qinchuan cattle (Bos taurus). Molecular and Cellular Probes, 2015, 29, 167-171.	2.1	28