

Jianghong Wu

List of Publications by Year in descending order

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Version: 2024-02-01

15
papers

136
citations

1684188

5
h-index

1281871

11
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18
all docs

18
docs citations

18
times ranked

191
citing authors

#	ARTICLE	IF	CITATIONS
1	Comparative Analysis of the Microbiota Between Rumen and Duodenum of Twin Lambs Based on Diets of Ceratoides or Alfalfa. Polish Journal of Microbiology, 2021, 70, 175-187.	1.7	2
2	The Dynamic Change of Gene-Regulated Networks in Cashmere Goat Skin with Seasonal Variation. Biochemical Genetics, 2021, , 1.	1.7	2
3	Editorial: Multi-Omics Approaches to Study Complex Traits in Domestic Animals. Frontiers in Systems Biology, 2021, 1, .	0.7	1
4	Multiple omics analysis reveals that high fiber diets promote gluconeogenesis and inhibit glycolysis in muscle. BMC Genomics, 2020, 21, 660.	2.8	10
5	Physiological responses and transcriptome analysis of the Kochia prostrata (L.) Schrad. to seedling drought stress. AIMS Genetics, 2019, 06, 017-035.	1.9	3
6	Evolution and Expression of S100A7 Gene in Vertebrates. Biochemical Genetics, 2019, 57, 371-381.	1.7	4
7	Skin transcriptome reveals the dynamic changes in the Wnt pathway during integument morphogenesis of chick embryos. PLoS ONE, 2018, 13, e0190933.	2.5	20
8	Transcriptome Reveals That Genic SNPs Contributes to Heterosis in Cattle. , 2018, , .		1
9	Analyzing the miRNA-Gene Networks to Mine the Important miRNAs under Skin of Human and Mouse. BioMed Research International, 2016, 2016, 1-9.	1.9	5
10	Dissecting the biological relationship between TCGA miRNA and mRNA sequencing data using MMiRNA-Viewer. BMC Bioinformatics, 2016, 17, 336.	2.6	10
11	Single Base-Resolution Methylome of the Dizygotic Sheep. PLoS ONE, 2015, 10, e0142034.	2.5	5
12	Adaptive Evolution of the STRA6 Genes in Mammalian. PLoS ONE, 2014, 9, e108388.	2.5	6
13	Adaptive evolution of Hoxc13 genes in the origin and diversification of the vertebrate integument. , 2013, 320, n/a-n/a.		5
14	A Subset of Skin-Expressed microRNAs with Possible Roles in Goat and Sheep Hair Growth Based on Expression Profiling of Mammalian microRNAs. OMICS A Journal of Integrative Biology, 2007, 11, 385-396.	2.0	61
15	The impact of different feeds on DNA methylation, glycolysis/gluconeogenesis signaling pathway, and gene expression of sheep muscle. PeerJ, 0, 10, e13455.	2.0	0