

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
g-index

18
all docs

18
docs citations

18
times ranked

191
citing authors

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