

Bugao Li

List of Publications by Year in descending order

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Version: 2024-02-01

16
papers

144
citations

1307594

7
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1199594

12
g-index

18
all docs

18
docs citations

18
times ranked

187
citing authors

#	ARTICLE	IF	CITATIONS
1	Glutamic-oxaloacetic transaminase 1 regulates adipocyte differentiation by altering nicotinamide adenine dinucleotide phosphate content. <i>Animal Bioscience</i> , 2022, 35, 155-165.	2.0	2
2	Long non-coding RNAs in <i>Sus scrofa</i> ; ileum under starvation stress. <i>Animal Bioscience</i> , 2022, , .	2.0	0
3	miR-128 regulated the proliferation and autophagy in porcine adipose-derived stem cells through targeting the JNK signaling pathway. <i>Journal of Receptor and Signal Transduction Research</i> , 2021, 41, 196-201.	2.5	3
4	Mining Unknown Porcine Protein Isoforms by Tissue-based Map of Proteome Enhances Pig Genome Annotation. <i>Genomics, Proteomics and Bioinformatics</i> , 2021, 19, 772-786.	6.9	2
5	PRE-1 Revealed Previous Unknown Introgression Events in Eurasian Boars during the Middle Pleistocene. <i>Genome Biology and Evolution</i> , 2020, 12, 1751-1764.	2.5	9
6	Comparative Transcriptome Analyses of <i>Longissimus thoracis</i> Between Pig Breeds Differing in Muscle Characteristics. <i>Frontiers in Genetics</i> , 2020, 11, 526309.	2.3	15
7	<i>MyHC</i> 's developmental expression patterns and its effect on muscle fibre characteristics in pig. <i>Journal of Applied Animal Research</i> , 2020, 48, 176-183.	1.2	4
8	Novel alternatively spliced isoforms of pig ZNF280D and their diverse mRNA expression patterns. <i>Biologia (Poland)</i> , 2019, 74, 1699-1709.	1.5	0
9	Comparison of carcass traits, meat quality and expressions of <i>MyHCs</i> in muscles between Mashen and Large White pigs. <i>Italian Journal of Animal Science</i> , 2019, 18, 1410-1418.	1.9	18
10	A comparison of dynamic distributions of intestinal microbiota between Large White and Chinese Shanxi Black pigs. <i>Archives of Microbiology</i> , 2019, 201, 357-367.	2.2	14
11	Identification of candidate genes of growth traits in pigs using RNA-sequencing. <i>Italian Journal of Animal Science</i> , 2019, 18, 279-286.	1.9	4
12	Selection of candidate genes affecting meat quality and preliminary exploration of related molecular mechanisms in the Mashen pig. <i>Asian-Australasian Journal of Animal Sciences</i> , 2019, 32, 1084-1094.	2.4	22
13	Profiling long noncoding RNA of multi-tissue transcriptome enhances porcine noncoding genome annotation. <i>Epigenomics</i> , 2018, 10, 301-320.	2.1	22
14	Factors affecting GEBV accuracy with single-step Bayesian models. <i>Heredity</i> , 2018, 120, 100-109.	2.6	10
15	Novel splice isoforms of pig myoneurin and their diverse mRNA expression patterns. <i>Asian-Australasian Journal of Animal Sciences</i> , 2018, 31, 1581-1590.	2.4	6
16	Study on the Developmental Expression of <i>Lbx1</i> Gene in <i>Longissimus Dorsi</i> of Mashen and Large White Pigs. <i>Italian Journal of Animal Science</i> , 2015, 14, 3720.	1.9	13