

Jiying Wang

List of Publications by Year in descending order

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Version: 2024-02-01

15
papers

2,632
citations

1163117

8
h-index

1058476

14
g-index

16
all docs

16
docs citations

16
times ranked

3584
citing authors

#	ARTICLE	IF	CITATIONS
1	Novel p19 Protein Engages IL-12p40 to Form a Cytokine, IL-23, with Biological Activities Similar as Well as Distinct from IL-12. <i>Immunity</i> , 2000, 13, 715-725.	14.3	2,463
2	Genome-wide detection of selection signatures in Chinese indigenous Laiwu pigs revealed candidate genes regulating fat deposition in muscle. <i>BMC Genetics</i> , 2018, 19, 31.	2.7	40
3	Muscle Transcriptome Analysis Reveals Potential Candidate Genes and Pathways Affecting Intramuscular Fat Content in Pigs. <i>Frontiers in Genetics</i> , 2020, 11, 877.	2.3	28
4	Genome-wide association study for intramuscular fat content in Chinese Lulai black pigs. <i>Asian-Australasian Journal of Animal Sciences</i> , 2019, 32, 607-613.	2.4	22
5	Selection of Reference Genes for Gene Expression Studies in Porcine Whole Blood and Peripheral Blood Mononuclear Cells under Polyinosinic:Polycytidylic Acid Stimulation. <i>Asian-Australasian Journal of Animal Sciences</i> , 2014, 27, 471-478.	2.4	17
6	A comparison of methods for effective differentiation of the frozen-thawed 3T3-L1 cells. <i>Analytical Biochemistry</i> , 2019, 568, 57-64.	2.4	16
7	Transcriptomic Analysis Identifies Candidate Genes and Gene Sets Controlling the Response of Porcine Peripheral Blood Mononuclear Cells to Poly I:C Stimulation. <i>G3: Genes, Genomes, Genetics</i> , 2016, 6, 1267-1275.	1.8	11
8	Comparative gene expression profiling of muscle reveals potential candidate genes affecting drip loss in pork. <i>BMC Genetics</i> , 2019, 20, 89.	2.7	10
9	Identification of reference microRNAs for quantitative expression analysis in porcine peripheral blood mononuclear cells treated with polyinosinic-polycytidylic acid. <i>International Journal of Immunogenetics</i> , 2015, 42, 217-225.	1.8	5
10	MicroRNA Transcriptome of Poly I:C-Stimulated Peripheral Blood Mononuclear Cells Reveals Evidence for MicroRNAs in Regulating Host Response to RNA Viruses in Pigs. <i>International Journal of Molecular Sciences</i> , 2016, 17, 1601.	4.1	5
11	Weighted gene co-expression network analysis reveals potential candidate genes affecting drip loss in pork. <i>Animal Genetics</i> , 2020, 51, 855-865.	1.7	5
12	High-density single nucleotide polymorphism chip-based conservation genetic analysis of indigenous pig breeds from Shandong Province, China. <i>Animal Bioscience</i> , 2021, 34, 1123-1133.	2.0	4
13	Association of the Porcine Cluster of Differentiation 4 Gene with T Lymphocyte Subpopulations and Its Expression in Immune Tissues. <i>Asian-Australasian Journal of Animal Sciences</i> , 2013, 26, 463-469.	2.4	3
14	Correlation analyses of CpG island methylation of cluster of differentiation 4 protein with gene expression and T lymphocyte subpopulation traits. <i>Asian-Australasian Journal of Animal Sciences</i> , 2018, 31, 1141-1149.	2.4	2
15	Differentiation proliferative capacity of skeletal muscle satellite cells from Dapulian and Landrace pigs. <i>Italian Journal of Animal Science</i> , 2020, 19, 574-585.	1.9	0