

Jiabo Wang

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3058973/publications.pdf>

Version: 2024-02-01

10
papers

927
citations

1651377

6
h-index

1637695

9
g-index

15
all docs

15
docs citations

15
times ranked

1175
citing authors

#	ARTICLE	IF	CITATIONS
1	Comparative transcriptome analysis of winter yaks in plateau and plain. <i>Reproduction in Domestic Animals</i> , 2022, 57, 64-71.	0.6	2
2	Identification of genetic loci associated with rough dwarf disease resistance in maize by integrating GWAS and linkage mapping. <i>Plant Science</i> , 2022, 315, 111100.	1.7	12
3	Performing Genome-Wide Association Studies with Multiple Models Using GAPIT. <i>Methods in Molecular Biology</i> , 2022, , 199-217.	0.4	1
4	Genome-Wide DNA Methylation and Hydroxymethylation Changes Revealed Epigenetic Regulation of Neuromodulation and Myelination in Yak Hypothalamus. <i>Frontiers in Genetics</i> , 2021, 12, 592135.	1.1	2
5	GAPIT Version 3: Boosting Power and Accuracy for Genomic Association and Prediction. <i>Genomics, Proteomics and Bioinformatics</i> , 2021, 19, 629-640.	3.0	345
6	Single Nucleotide Polymorphism Detection for Peach Gummosis Disease Resistance by Genome-Wide Association Study. <i>Frontiers in Plant Science</i> , 2021, 12, 763618.	1.7	4
7	The whole-transcriptome landscape of muscle and adipose tissues reveals the ceRNA regulation network related to intramuscular fat deposition in yak. <i>BMC Genomics</i> , 2020, 21, 347.	1.2	40
8	Detection and integrated analysis of lncRNA and mRNA relevant to plateau adaptation of Yak. <i>Reproduction in Domestic Animals</i> , 2020, 55, 1461-1469.	0.6	8
9	Expanding the BLUP alphabet for genomic prediction adaptable to the genetic architectures of complex traits. <i>Heredity</i> , 2018, 121, 648-662.	1.2	53
10	GAPIT Version 2: An Enhanced Integrated Tool for Genomic Association and Prediction. <i>Plant Genome</i> , 2016, 9, plantgenome2015.11.0120.	1.6	421