Ping Yan

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/4480749/publications.pdf

Version: 2024-02-01

19 papers	146 citations	1307594 7 h-index	1281871 11 g-index
19 all docs	19 docs citations	19 times ranked	117 citing authors

#	Article	IF	Citations
1	Differential proteomic analysis demonstrates follicle fluid participate immune reaction and protein translation in yak. BMC Veterinary Research, 2022, 18, 34.	1.9	3
2	Copy number variation (CNV) of the AHR gene in the Ashidan yak and its association with growth traits. Gene, 2022, 826, 146454.	2.2	1
3	The transcriptome-wide N6-methyladenosine (m6A) map profiling reveals the regulatory role of m6A in the yak ovary. BMC Genomics, 2022, 23, 358.	2.8	7
4	Secondary Structural Transformation of Bovine Lactoferricin Affects Its Antibacterial Activity. Probiotics and Antimicrobial Proteins, 2021, 13, 873-884.	3.9	5
5	Copy number variation of the HPGDS gene in the Ashidan yak and its associations with growth traits. Gene, 2021, 772, 145382.	2.2	11
6	Mitogenomic diversity and phylogeny analysis of yak (Bos grunniens). BMC Genomics, 2021, 22, 325.	2.8	18
7	The Study on Potential Biomarker in Rat After Withdrawal of Cimaterol Using Untargeted Metabonomics. Chromatographia, 2021, 84, 677-686.	1.3	2
8	Using weighted gene co-expression network analysis (WGCNA) to identify the hub genes related to hypoxic adaptation in yak (Bos grunniens). Genes and Genomics, 2021, 43, 1231-1246.	1.4	12
9	Fat deposition in yak during different phenological seasons. Livestock Science, 2021, 251, 104671.	1.6	3
10	Effect of intramolecular disulfide bond of bovine lactoferricin on its molecular structure and antibacterial activity against Trueperella pyogenes separated from cow milk with mastitis. BMC Veterinary Research, 2020, 16, 401.	1.9	2
11	Multi-residue Determination of Bisphenol Compounds in Feed Using Ultrasound-Assisted Extraction and Dispersive Solid-Phase Extract Followed by High-Performance Liquid Chromatography with Fluorescence Detector. Chromatographia, 2020, 83, 1423-1433.	1.3	6
12	Reference gene selection and myosin heavy chain (MyHC) isoform expression in muscle tissues of domestic yak (Bos grunniens). PLoS ONE, 2020, 15, e0228493.	2.5	11
13	Analysis of Hematological Traits in Polled Yak by Genome-Wide Association Studies Using Individual SNPs and Haplotypes. Genes, 2019, 10, 463.	2.4	17
14	Dietary Energy Levels Affect Growth Performance through Growth Hormone and Insulin-Like Growth Factor 1 in Yak (Bos grunniens). Animals, 2019, 9, 39.	2.3	13
15	Characterisation of the complete mitochondrial genome of the Jinchuan Yak (Bos grunniens). Mitochondrial DNA Part B: Resources, 2019, 4, 3856-3857.	0.4	0
16	The Selection of Reference Genes for Quantitative Real-Time PCR in the Ashidan Yak Mammary Gland During Lactation and Dry Period. Animals, 2019, 9, 943.	2.3	22
17	Population genetic variations of the matrix metalloproteinases-3 gene revealed hypoxia adaptation in domesticated yaks (Bos grunniens). Asian-Australasian Journal of Animal Sciences, 2019, 32, 1801-1808.	2.4	2
18	The complete mitochondrial genome of Hequ Tibetan Mastiff Canis lupus familiaris (Carnivora:) Tj ETQq0 0 0 rgBT /Overlock 10 T	0.7	0

#	Article	IF	Citations
19	The complete mitochondrial genome of the Qinghai Plateau yak Bos grunniens (Cetartiodactyla:) Tj ETQq1 1 0.784314 rgBT /Ove	0.7	11