

Xingang Feng

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9122406/publications.pdf>

Version: 2024-02-01

11
papers

115
citations

1478505

6
h-index

1372567

10
g-index

11
all docs

11
docs citations

11
times ranked

133
citing authors

#	ARTICLE	IF	CITATIONS
1	Analyses of mitogenomic markers shed light on the divergence, population dynamics, and demographic history of Pakistani chickens. <i>Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis</i> , 2021, 32, 34-42.	0.7	0
2	Suppression of hyaluronidase reduces invasion and establishment of <i>Haemonchus contortus</i> larvae in sheep. <i>Veterinary Research</i> , 2020, 51, 106.	3.0	1
3	A Whole Genome Re-Sequencing Based GWA Analysis Reveals Candidate Genes Associated with Ivermectin Resistance in <i>Haemonchus contortus</i> . <i>Genes</i> , 2020, 11, 367.	2.4	26
4	Analysis of genome-wide SNPs based on 2b-RAD sequencing of pooled samples reveals signature of selection in different populations of <i>Haemonchus contortus</i> . <i>Journal of Biosciences</i> , 2019, 44, 1.	1.1	9
5	Microsatellite analysis reveals extensive gene flow, and lack of population structure in the farm populations of <i>Haemonchus contortus</i> in northern China. <i>Parasitology International</i> , 2019, 73, 101959.	1.3	2
6	Ethnogenetic analysis reveals that Kohistanis of Pakistan were genetically linked to west Eurasians by a probable ancestral genepool from Eurasian steppe in the bronze age. <i>Mitochondrion</i> , 2019, 47, 82-93.	3.4	2
7	Population structure, genetic diversity and phylogenetic analysis of different rural and commercial chickens of Pakistan using complete sequence of mtDNA D-loop. <i>Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis</i> , 2019, 30, 273-280.	0.7	9
8	Analysis of genome-wide SNPs based on 2b-RAD sequencing of pooled samples reveals signature of selection in different populations of. <i>Journal of Biosciences</i> , 2019, 44, .	1.1	5
9	Genome-wide SNP analysis using 2b-RAD sequencing identifies the candidate genes putatively associated with resistance to ivermectin in <i>Haemonchus contortus</i> . <i>Parasites and Vectors</i> , 2017, 10, 31.	2.5	47
10	Microarray Analysis of Gene Expression Profiles of <i>Schistosoma japonicum</i> Derived from Less-Susceptible Host Water Buffalo and Susceptible Host Goat. <i>PLoS ONE</i> , 2013, 8, e70367.	2.5	8
11	Characterization and expression of a novel Frizzled 9 gene in <i>Schistosoma japonicum</i> . <i>Gene Expression Patterns</i> , 2011, 11, 263-270.	0.8	6