

Larissa Fernanda Simielli Fonseca

List of Publications by Year in descending order

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Version: 2024-02-01

17
papers

297
citations

933447

10
h-index

940533

16
g-index

17
all docs

17
docs citations

17
times ranked

456
citing authors

#	ARTICLE	IF	CITATIONS
1	Genes and proteins associated with ribeye area and meat tenderness in a commercial Nelore cattle population. <i>Genome</i> , 2022, 65, 229-240.	2.0	2
2	Small genetic variation affecting mRNA isoforms associated with marbling and meat color in beef cattle. <i>Functional and Integrative Genomics</i> , 2022, , 1.	3.5	0
3	Gene expression profiling and identification of hub genes in Nelore cattle with different marbling score levels. <i>Genomics</i> , 2020, 112, 873-879.	2.9	31
4	Transcriptome profiling of muscle in Nelore cattle phenotypically divergent for the ribeye muscle area. <i>Genomics</i> , 2020, 112, 1257-1263.	2.9	13
5	Use of gene expression profile to identify potentially relevant transcripts to myofibrillar fragmentation index trait. <i>Functional and Integrative Genomics</i> , 2020, 20, 609-619.	3.5	10
6	Prediction of hub genes associated with intramuscular fat content in Nelore cattle. <i>BMC Genomics</i> , 2019, 20, 520.	2.8	24
7	Influence of X-chromosome markers on reproductive traits of beef cattle. <i>Livestock Science</i> , 2019, 220, 152-157.	1.6	5
8	Expression profile of the CSF3 and LPO genes in milk from buffalo (<i>Bubalus bubalis</i>) with and without mastitis. <i>Molecular and Cellular Probes</i> , 2018, 41, 39-42.	2.1	1
9	Genomic association for sexual precocity in beef heifers using pre-selection of genes and haplotype reconstruction. <i>PLoS ONE</i> , 2018, 13, e0190197.	2.5	20
10	Differences in global gene expression in muscle tissue of Nelore cattle with divergent meat tenderness. <i>BMC Genomics</i> , 2017, 18, 945.	2.8	32
11	Detection of differentially methylated regions of irradiated fig tree selections. <i>Scientia Agricola</i> , 2017, 74, 285-293.	1.2	2
12	Gene expression profile of intramuscular muscle in Nelore cattle with extreme values of fatty acid. <i>BMC Genomics</i> , 2016, 17, 972.	2.8	49
13	Expression of genes related to mitochondrial function in Nelore cattle divergently ranked on residual feed intake. <i>Molecular Biology Reports</i> , 2015, 42, 559-565.	2.3	20
14	Differential gene expression in drought-tolerant sugarcane roots. <i>Genetics and Molecular Research</i> , 2015, 14, 7196-7207.	0.2	34
15	Polymorphism analysis in genes of the somatotropic axis in Nelore cattle selected for growth. <i>Gene</i> , 2014, 545, 215-219.	2.2	9
16	Evaluation of TFAM and FABP4 gene polymorphisms in three lines of Nelore cattle selected for growth. <i>Genetics and Molecular Research</i> , 2010, 9, 2050-2059.	0.2	14
17	Assessment of DGAT1 and LEP gene polymorphisms in three Nelore (<i>Bos indicus</i>) lines selected for growth and their relationship with growth and carcass traits ¹ . <i>Journal of Animal Science</i> , 2010, 88, 435-441.	0.5	31