

Warren M Snelling

List of Publications by Year in descending order

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Version: 2024-02-01

22
papers

2,156
citations

759233

12
h-index

713466

21
g-index

23
all docs

23
docs citations

23
times ranked

2820
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|------|-----------|
| 1 | Genetic parameters, heterosis, and breed effects for body condition score and mature cow weight in beef cattle. <i>Journal of Animal Science</i> , 2022, 100, . | 0.5 | 6 |
| 2 | Influence of environmental factors and genetic variation on mitochondrial DNA copy number. <i>Journal of Animal Science</i> , 2022, 100, . | 0.5 | 6 |
| 3 | Breeding Sustainable Beef Cows: Reducing Weight and Increasing Productivity. <i>Animals</i> , 2022, 12, 1745. | 2.3 | 1 |
| 4 | Genetic changes in beef cow traits following selection for calving ease. <i>Translational Animal Science</i> , 2021, 5, txab009. | 1.1 | 1 |
| 5 | Using Genomics to Measure Phenomics: Repeatability of Bull Prolificacy in Multiple-Bull Pastures. <i>Agriculture (Switzerland)</i> , 2021, 11, 603. | 3.1 | 2 |
| 6 | Breed and heterotic effects for mature weight in beef cattle. <i>Journal of Animal Science</i> , 2021, 99, . | 0.5 | 8 |
| 7 | Assessment of Imputation from Low-Pass Sequencing to Predict Merit of Beef Steers. <i>Genes</i> , 2020, 11, 1312. | 2.4 | 55 |
| 8 | Heritability and genetic correlations of feed intake, body weight gain, residual gain, and residual feed intake of beef cattle as heifers and cows. <i>Journal of Animal Science</i> , 2020, 98, . | 0.5 | 34 |
| 9 | De novo assembly of the cattle reference genome with single-molecule sequencing. <i>GigaScience</i> , 2020, 9, . | 6.4 | 380 |
| 10 | Comparison of different functions to describe growth from weaning to maturity in crossbred beef cattle1. <i>Journal of Animal Science</i> , 2019, 97, 1523-1533. | 0.5 | 10 |
| 11 | Genetic correlations among weight and cumulative productivity of crossbred beef cows1. <i>Journal of Animal Science</i> , 2019, 97, 63-77. | 0.5 | 15 |
| 12 | Reducing the period of data collection for intake and gain to improve response to selection for feed efficiency in beef cattle. <i>Journal of Animal Science</i> , 2018, 96, 854-866. | 0.5 | 8 |
| 13 | RNA-Seq Meta-analysis identifies genes in skeletal muscle associated with gain and intake across a multi-season study of crossbred beef steers. <i>BMC Genomics</i> , 2018, 19, 430. | 2.8 | 21 |
| 14 | Profile of the Spleen Transcriptome in Beef Steers with Variation in Gain and Feed Intake. <i>Frontiers in Genetics</i> , 2016, 7, 127. | 2.3 | 14 |
| 15 | Transcriptome differences in the rumen of beef steers with variation in feed intake and gain. <i>Gene</i> , 2016, 586, 12-26. | 2.2 | 45 |
| 16 | Multi-Tissue Omics Analyses Reveal Molecular Regulatory Networks for Puberty in Composite Beef Cattle. <i>PLoS ONE</i> , 2014, 9, e102551. | 2.5 | 125 |
| 17 | A multiway analysis for identifying high integrity bovine BACs. <i>BMC Genomics</i> , 2009, 10, 46. | 2.8 | 4 |
| 18 | The Genome Sequence of Taurine Cattle: A Window to Ruminant Biology and Evolution. <i>Science</i> , 2009, 324, 522-528. | 12.6 | 1,038 |

| # | ARTICLE | IF | CITATIONS |
|----|---|-----|-----------|
| 19 | A physical map of the bovine genome. <i>Genome Biology</i> , 2007, 8, R165. | 9.6 | 73 |
| 20 | Linkage mapping bovine EST-based SNP. <i>BMC Genomics</i> , 2005, 6, 74. | 2.8 | 58 |
| 21 | A Comprehensive Genetic Map of the Cattle Genome Based on 3802 Microsatellites. <i>Genome Research</i> , 2004, 14, 1987-1998. | 5.5 | 237 |
| 22 | Integrating linkage and radiation hybrid mapping data for bovine chromosome 15. <i>BMC Genomics</i> , 2004, 5, 77. | 2.8 | 14 |