Jennifer M Thomson

List of Publications by Year in descending order

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623734 526287 36 938 14 27 citations g-index h-index papers 37 37 37 1381 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	Average kinship within bighorn sheep populations is associated with connectivity, augmentation, and bottlenecks. Ecosphere, 2022, 13 , .	2.2	3
2	Fat Deposition and Fat Effects on Meat Quality—A Review. Animals, 2022, 12, 1550.	2.3	25
3	Identification of Genomic Regions for Carcass Quality Traits within the American Simmental Association Carcass Merit Program. Animals, 2021, 11, 471.	2.3	1
4	1H NMR based metabolic profiling distinguishes the differential impact of capture techniques on wild bighorn sheep. Scientific Reports, 2021, 11, 11308.	3.3	3
5	Gene Expression and Carcass Traits Are Different between Different Quality Grade Groups in Red-Faced Hereford Steers. Animals, 2021, 11, 1910.	2.3	5
6	Differential haptoglobin responsiveness to a <i>Mannheimia haemolytica</i> challenge altered immunologic, physiologic, and behavior responses in beef steers. Journal of Animal Science, 2021, 99, .	0.5	2
7	A comparative approach to refine molecular mechanisms impacting meat quality and carcass characteristics. Translational Animal Science, 2021, 5, S189-S194.	1.1	0
8	Evaluating wildlife translocations using genomics: A bighorn sheep case study. Ecology and Evolution, 2020, 10, 13687-13704.	1.9	16
9	Phenotypic and genetic differences in Rambouillet lines divergently selected for reproductive rate over 50 years 1,2. Translational Animal Science, 2020, 4, S90-S93.	1.1	1
10	277 Phenotypic and genetic differences in Rambouillet lines divergently selected for reproductive rate over 50 years. Journal of Animal Science, 2020, 98, 206-206.	0.5	0
11	44 Inbreeding levels of the Line 1 and Line 4 Hereford cattle populations. Journal of Animal Science, 2020, 98, 18-18.	0.5	O
12	PSXI-19 Investigating the Relationship Between Temperament and Performance Traits in Feedlot Cattle. Journal of Animal Science, 2020, 98, 383-384.	0.5	0
13	Blood lactate and rectal temperature can predict exit velocity of beef feedlot steers. Translational Animal Science, 2019, 3, 1530-1542.	1.1	8
14	PSVIII-24 Inbreeding levels of the Line 4 Hereford cattle population. Journal of Animal Science, 2019, 97, 269-269.	0.5	0
15	Gene expression in muscle and adipose tissue of steers classed as Choice or Standard. , 2019, , .		0
16	Identification of genetic markers and QTL for carcass quality traits within the American Simmental Association Carcass Merit Program1. Translational Animal Science, 2018, 2, S39-S43.	1.1	0
17	Genetic Markers Are Associated with the Ruminal Microbiome and Metabolome in Grain and Sugar Challenged Dairy Heifers. Frontiers in Genetics, 2018, 9, 62.	2.3	24
18	Evaluating sample size to estimate genetic management metrics in the genomics era. Molecular Ecology Resources, 2018, 18, 1077-1091.	4.8	27

#	Article	IF	CITATIONS
19	Feed efficiency phenotypes in lambs involve changes in ruminal, colonic, and small-intestine-located microbiota1. Journal of Animal Science, 2017, 95, 2585-2592.	0.5	92
20	Feed efficiency phenotypes in lambs involve changes in ruminal, colonic, and small-intestine-located microbiota. Journal of Animal Science, 2017, 95, 2585.	0.5	42
21	0777 Repeatability of residual feed intake and indices of body composition in growing Columbia ewes fed the same diet. Journal of Animal Science, 2016, 94, 373-374.	0.5	2
22	095 Metabolomic profiling for identification of biomarkers associated with temperament in feedlot cattle. Journal of Animal Science, 2016, 94, 47-47.	0.5	0
23	Impacts of environment on gene expression and epigenetic modification in grazing animals. Journal of Animal Science, 2016, 94, 63-73.	0.5	1
24	Invited review: Recommendations for reporting intervention studies on reproductive performance in dairy cattle: Improving design, analysis, and interpretation of research on reproduction. Journal of Dairy Science, 2016, 99, 1-17.	3.4	85
25	The use of blood lactate concentration as an indicator of temperament and its impact on growth rate and tenderness of steaks from Simmental×Angus steers. Meat Science, 2015, 103, 68-74.	5.5	13
26	Characterization of the Vaginal Microbiota of Ewes and Cows Reveals a Unique Microbiota with Low Levels of Lactobacilli and Near-Neutral pH. Frontiers in Veterinary Science, 2014, 1, 19.	2.2	108
27	Plasma metabolites associated with residual feed intake and other productivity performance traits in beef cattle. Livestock Science, 2014, 165, 200-211.	1.6	63
28	Candidate genes and biological pathways associated with carcass quality traits in beef cattle. Canadian Journal of Animal Science, 2013, 93, 295-306.	1.5	12
29	Candidate genes and single nucleotide polymorphisms associated with variation in residual feed intake in beef cattle1. Journal of Animal Science, 2013, 91, 3502-3513.	0.5	44
30	Characterization of the longissimus lumborum transcriptome response to adding propionate to the diet of growing Angus beef steers. Physiological Genomics, 2012, 44, 543-550.	2.3	20
31	The identification of candidate genes and SNP markers for classical bovine spongiform encephalopathy susceptibility. Prion, 2012, 6, 461-469.	1.8	3
32	Differential expression of genes in adipose tissue of first-lactation dairy cattle. Journal of Dairy Science, 2011, 94, 361-369.	3.4	51
33	Whole genome resequencing of black Angus and Holstein cattle for SNP and CNV discovery. BMC Genomics, 2011, 12, 559.	2.8	153
34	The effect of Ala293Val single nucleotide polymorphism in the stearoyl-CoA desaturase gene on conjugated linoleic acid concentration in milk fat of dairy cows. Canadian Journal of Animal Science, 2010, 90, 575-584.	1.5	2
35	Effects of Chromium Propionate on Response to an Intravenous Glucose Tolerance Test in Growing Holstein Heifers. Journal of Dairy Science, 2007, 90, 3467-3474.	3.4	52
36	Expression of Lipolytic Genes in the Adipose Tissue of Pregnant and Lactating Holstein Dairy Cattle. Journal of Dairy Science, 2007, 90, 5237-5246.	3.4	80