Juan J Loor

List of Publications by Citations

440 papers	12,278 citations	55 h-index	90 g-index
484 ext. papers	15,394 ext. citations	3.7 avg, IF	6.75 L-index

#	Paper	IF	Citations
440	Gene networks driving bovine milk fat synthesis during the lactation cycle. *BMC Genomics*, **2008**, 9, 366	4.5	499
439	Regulation of energy metabolism by long-chain fatty acids. *Progress in Lipid Research*, **2014**, 53, 124-44	14.3	369
438	Identification of a missense mutation in the bovine ABCG2 gene with a major effect on the QTL on chromosome 6 affecting milk yield and composition in Holstein cattle. *Genome Research*, **2005**, 15, 936-44	9.7	273
437	Nutrition-induced ketosis alters metabolic and signaling gene networks in liver of periparturient dairy cows. *Physiological Genomics*, **2007**, 32, 105-16	3.6	243
436	Identification of reference genes for quantitative real-time PCR in the bovine mammary gland during the lactation cycle. *Physiological Genomics*, **2007**, 29, 312-9	3.6	237
435	Relationship among trans and conjugated fatty acids and bovine milk fat yield due to dietary concentrate and linseed oil. *Journal of Dairy Science*, **2005**, 88, 726-40	4	223
434	Peroxisome proliferator-activated receptor-gamma activation and long-chain fatty acids alter lipogenic gene networks in bovine mammary epithelial cells to various extents. *Journal of Dairy Science*, **2009**, 92, 4276-89	4	213
433	Biohydrogenation, duodenal flow, and intestinal digestibility of trans fatty acids and conjugated linoleic acids in response to dietary forage:concentrate ratio and linseed oil in dairy cows. *Journal of Dairy Science*, **2004**, 87, 2472-85	4	209
432	Gene networks driving bovine mammary protein synthesis during the lactation cycle. *Bioinformatics and Biology Insights*, **2011**, 5, 83-98	5.3	193
431	Temporal gene expression profiling of liver from periparturient dairy cows reveals complex adaptive mechanisms in hepatic function. *Physiological Genomics*, **2005**, 23, 217-26	3.6	173
430	Physiological and pathological adaptations in dairy cows that may increase susceptibility to periparturient diseases and disorders. *Italian Journal of Animal Science*, **2005**, 4, 323-344	2.2	168
429	ACSL1, AGPAT6, FABP3, LPIN1, and SLC27A6 are the most abundant isoforms in bovine mammary tissue and their expression is affected by stage of lactation. *Journal of Nutrition*, **2008**, 138, 1019-24	4.1	157
428	Plane of nutrition prepartum alters hepatic gene expression and function in dairy cows as assessed by longitudinal transcript and metabolic profiling. *Physiological Genomics*, **2006**, 27, 29-41	3.6	150
427	Supplemental Smartamine M or MetaSmart during the transition period benefits postpartal cow performance and blood neutrophil function. *Journal of Dairy Science*, **2013**, 96, 6248-63	4	121
426	Effect of linseed oil supplementation on ruminal digestion in dairy cows fed diets with different forage:concentrate ratios. *Journal of Dairy Science*, **2003**, 86, 3999-4007	4	112
425	Functional Role of PPARs in Ruminants: Potential Targets for Fine-Tuning Metabolism during Growth and Lactation. *PPAR Research*, **2013**, 2013, 684159	4.3	107
424	High-throughput Methods Redefine the Rumen Microbiome and Its Relationship with Nutrition and Metabolism. *Bioinformatics and Biology Insights*, **2014**, 8, 109-25	5.3	105

423	Housekeeping gene expression in bovine liver is affected by physiological state, feed intake, and dietary treatment. *Journal of Dairy Science*, **2007**, 90, 2246-52	4	101
422	Biomarkers of inflammation, metabolism, and oxidative stress in blood, liver, and milk reveal a better immunometabolic status in peripartal cows supplemented with Smartamine M or MetaSmart. *Journal of Dairy Science*, **2014**, 97, 7437-50	4	100
421	Systems physiology in dairy cattle: nutritional genomics and beyond. *Annual Review of Animal Biosciences*, **2013**, 1, 365-92	13.7	96
420	High-concentrate diets and polyunsaturated oils alter trans and conjugated isomers in bovine rumen, blood, and milk. *Journal of Dairy Science*, **2005**, 88, 3986-99	4	96
419	Soybean oil and linseed oil supplementation affect profiles of ruminal microorganisms in dairy cows. *Animal*, **2009**, 3, 1562-9	3.1	94
418	Exogenous conjugated linoleic acid isomers reduce bovine milk fat concentration and yield by inhibiting de novo fatty acid synthesis. *Journal of Nutrition*, **1998**, 128, 2411-9	4.1	91
417	Reduced fatty acid synthesis and desaturation due to exogenous trans10,cis12-CLA in cows fed oleic or linoleic oil. *Journal of Dairy Science*, **2003**, 86, 1354-69	4	90
416	Adipogenic and energy metabolism gene networks in longissimus lumborum during rapid post-weaning growth in Angus and Angus x Simmental cattle fed high-starch or low-starch diets. *BMC Genomics*, **2009**, 10, 142	4.5	89
415	Old and new stories: revelations from functional analysis of the bovine mammary transcriptome during the lactation cycle. *PLoS ONE*, **2012**, 7, e33268	3.7	88
414	Genomics of metabolic adaptations in the peripartal cow. *Animal*, **2010**, 4, 1110-39	3.1	88
413	Transcriptome profiling of the small intestinal epithelium in germfree versus conventional piglets. *BMC Genomics*, **2007**, 8, 215	4.5	87
412	Intestinal flow and digestibility of trans fatty acids and conjugated linoleic acids (CLA) in dairy cows fed a high-concentrate diet supplemented with fish oil, linseed oil, or sunflower oil. *Animal Feed Science and Technology*, **2005**, 119, 203-225	3	82
411	Better postpartal performance in dairy cows supplemented with rumen-protected methionine compared with choline during the peripartal period. *Journal of Dairy Science*, **2016**, 99, 8716-8732	4	81
410	MicroRNA expression patterns in the bovine mammary gland are affected by stage of lactation. *Journal of Dairy Science*, **2012**, 95, 6529-35	4	80
409	Nutritional management of the transition cow in the 21st century – a paradigm shift in thinking. *Animal Production Science*, **2013**, 53, 1000	1.4	79
408	Blood immunometabolic indices and polymorphonuclear neutrophil function in peripartum dairy cows are altered by level of dietary energy prepartum. *Journal of Dairy Science*, **2012**, 95, 1749-58	4	79
407	52 Young Scholar Presentation: Maternal supply of methionine during late-pregnancy alters in utero and neonatal development, hepatic one-carbon metabolism,and innate immune response in Holstein calves. *Journal of Animal Science*, **2019**, 97, 26-27	0.7	78
406	186 Young Scholar Presentation: Immunometabolism during periods of negative nutrient balance or heat stress is altered by dietary methyl donor supply in dairy cows. *Journal of Animal Science*, **2020**, 98, 13-14	0.7	78

405	Gene network and pathway analysis of bovine mammary tissue challenged with Streptococcus uberis reveals induction of cell proliferation and inhibition of PPARgamma signaling as potential mechanism for the negative relationships between immune response and lipid metabolism. *BMC Genomics*, **2009**, 10, 542	4.5	78
404	Effects of ruminal or duodenal supply of fish oil on milk fat secretion and profiles of trans-fatty acids and conjugated linoleic acid isomers in dairy cows fed maize silage. *Animal Feed Science and Technology*, **2005**, 119, 227-246	3	77
403	Induction of Subacute Ruminal Acidosis Affects the Ruminal Microbiome and Epithelium. *Frontiers in Microbiology*, **2016**, 7, 701	5.7	77
402	Overfeeding a moderate energy diet prepartum does not impair bovine subcutaneous adipose tissue insulin signal transduction and induces marked changes in peripartal gene network expression. *Journal of Dairy Science*, **2012**, 95, 4333-51	4	76
401	Identification of internal control genes for quantitative polymerase chain reaction in mammary tissue of lactating cows receiving lipid supplements. *Journal of Dairy Science*, **2009**, 92, 2007-19	4	75
400	Liver lipid content and inflammometabolic indices in peripartal dairy cows are altered in response to prepartal energy intake and postpartal intramammary inflammatory challenge. *Journal of Dairy Science*, **2013**, 96, 918-35	4	74
399	Rumen-protected methionine compared with rumen-protected choline improves immunometabolic status in dairy cows during the peripartal period. *Journal of Dairy Science*, **2016**, 99, 8956-8969	4	72
398	Ethyl-cellulose rumen-protected methionine enhances performance during the periparturient period and early lactation in Holstein dairy cows. *Journal of Dairy Science*, **2017**, 100, 7455-7467	4	70
397	Dietary-induced negative energy balance has minimal effects on innate immunity during a Streptococcus uberis mastitis challenge in dairy cows during midlactation. *Journal of Dairy Science*, **2009**, 92, 4301-16	4	70
396	A novel dynamic impact approach (DIA) for functional analysis of time-course omics studies: validation using the bovine mammary transcriptome. *PLoS ONE*, **2012**, 7, e32455	3.7	69
395	Role of metabolic and cellular proliferation genes in ruminal development in response to enhanced plane of nutrition in neonatal Holstein calves. *Journal of Dairy Science*, **2012**, 95, 1807-20	4	67
394	Body condition score at calving affects systemic and hepatic transcriptome indicators of inflammation and nutrient metabolism in grazing dairy cows. *Journal of Dairy Science*, **2015**, 98, 1019-32	4	65
393	Fine metabolic regulation in ruminants via nutrient-gene interactions: saturated long-chain fatty acids increase expression of genes involved in lipid metabolism and immune response partly through PPAR-□activation. *British Journal of Nutrition*, **2012**, 107, 179-91	3.6	63
392	Milk fat depression induced by dietary marine algae in dairy ewes: persistency of milk fatty acid composition and animal performance responses. *Journal of Dairy Science*, **2013**, 96, 524-32	4	61
391	Bioinformatics analysis of microRNA and putative target genes in bovine mammary tissue infected with Streptococcus uberis. *Journal of Dairy Science*, **2012**, 95, 6397-408	4	60
390	In vitro culture and characterization of a mammary epithelial cell line from Chinese Holstein dairy cow. *PLoS ONE*, **2009**, 4, e7636	3.7	60
389	Ethyl-cellulose rumen-protected methionine alleviates inflammation and oxidative stress and improves neutrophil function during the periparturient period and early lactation in Holstein dairy cows. *Journal of Dairy Science*, **2018**, 101, 480-490	4	59
388	Effects of precalving body condition score and prepartum feeding level on production, reproduction, and health parameters in pasture-based transition dairy cows. *Journal of Dairy Science*, **2015**, 98, 7164-82	4	57

387	Review: Enhancing gastrointestinal health in dairy cows. *Animal*, **2018**, 12, s399-s418	3.1	56
386	Maternal consumption of organic trace minerals alters calf systemic and neutrophil mRNA and microRNA indicators of inflammation and oxidative stress. *Journal of Dairy Science*, **2015**, 98, 7717-29	4	55
385	Nutrient digestion, biohydrogenation, and fatty acid profiles in blood plasma and milk fat from lactating Holstein cows fed canola oil or canolamide. *Animal Feed Science and Technology*, **2002**, 97, 65-82	3	55
384	Greater expression of TLR2, TLR4, and IL6 due to negative energy balance is associated with lower expression of HLA-DRA and HLA-A in bovine blood neutrophils after intramammary mastitis challenge with Streptococcus uberis. *Functional and Integrative Genomics*, **2010**, 10, 53-61	3.8	54
383	Calving body condition score affects indicators of health in grazing dairy cows. *Journal of Dairy Science*, **2013**, 96, 5811-25	4	53
382	Smartamine M and MetaSmart supplementation during the peripartal period alter hepatic expression of gene networks in 1-carbon metabolism, inflammation, oxidative stress, and the growth hormone-insulin-like growth factor 1 axis pathways. *Journal of Dairy Science*, **2014**, 97, 7451-64	4	51
381	Acyl-CoA synthetase short-chain family member 2 (ACSS2) is regulated by SREBP-1 and plays a role in fatty acid synthesis in caprine mammary epithelial cells. *Journal of Cellular Physiology*, **2018**, 233, 1005-1016	7	50
380	Change in subcutaneous adipose tissue metabolism and gene network expression during the transition period in dairy cows, including differences due to sire genetic merit. *Journal of Dairy Science*, **2013**, 96, 2171-2182	4	50
379	Peroxisome proliferator-activated receptor-δ stimulates the synthesis of monounsaturated fatty acids in dairy goat mammary epithelial cells via the control of stearoyl-coenzyme A desaturase. *Journal of Dairy Science*, **2013**, 96, 7844-53	4	49
378	Assessing and managing body condition score for the prevention of metabolic disease in dairy cows. *Veterinary Clinics of North America - Food Animal Practice*, **2013**, 29, 323-36	4.6	49
377	A 7872 cDNA microarray and its use in bovine functional genomics. *Veterinary Immunology and Immunopathology*, **2005**, 105, 235-45	2	49
376	Biohydrogenation of unsaturated fatty acids in continuous culture fermenters during digestion of orchardgrass or red clover with three levels of ground corn supplementation. *Journal of Animal Science*, **2003**, 81, 1611-27	0.7	49
375	Characterization of 18:1 and 18:2 isomers produced during microbial biohydrogenation of unsaturated fatty acids from canola and soya bean oil in the rumen of lactating cows. *Journal of Animal Physiology and Animal Nutrition*, **2002**, 86, 422-32	2.6	49
374	Abundance of ruminal bacteria, epithelial gene expression, and systemic biomarkers of metabolism and inflammation are altered during the peripartal period in dairy cows. *Journal of Dairy Science*, **2015**, 98, 8940-51	4	48
373	Visceral adipose tissue mass in nonlactating dairy cows fed diets differing in energy density(1). *Journal of Dairy Science*, **2014**, 97, 3420-30	4	48
372	Trans18:1 and 18:2 isomers in blood plasma and milk fat of grazing cows fed a grain supplement containing solvent-extracted or mechanically extracted soybean meal. *Journal of Dairy Science*, **2002**, 85, 1197-207	4	47
371	Maternal rumen-protected methionine supplementation and its effect on blood and liver biomarkers of energy metabolism, inflammation, and oxidative stress in neonatal Holstein calves. *Journal of Dairy Science*, **2016**, 99, 6753-6763	4	47
370	miR-148a and miR-17-5p synergistically regulate milk TAG synthesis via PPARGC1A and PPARA in goat mammary epithelial cells. *RNA Biology*, **2017**, 14, 326-338	4.8	46

369	RNA-Seq reveals 10 novel promising candidate genes affecting milk protein concentration in the Chinese Holstein population. *Scientific Reports*, **2016**, 6, 26813	4.9	45
368	Circulating amino acids in blood plasma during the peripartal period in dairy cows with different liver functionality index. *Journal of Dairy Science*, **2016**, 99, 2257-2267	4	45
367	MiR-145 Regulates Lipogenesis in Goat Mammary Cells Via Targeting INSIG1 and Epigenetic Regulation of Lipid-Related Genes. *Journal of Cellular Physiology*, **2017**, 232, 1030-1040	7	45
366	Trans10,cis12-18:2 is a more potent inhibitor of de novo fatty acid synthesis and desaturation than cis9,trans11-18:2 in the mammary gland of lactating mice. *Journal of Nutrition*, **2004**, 134, 1362-8	4.1	45
365	Functional welfare using biochemical and molecular technologies to understand better the welfare state of peripartal dairy cattle. *Animal Production Science*, **2013**, 53, 931	1.4	44
364	iTRAQ-proteomics and bioinformatics analyses of mammary tissue from cows with clinical mastitis due to natural infection with Staphylococci aureus. *BMC Genomics*, **2014**, 15, 839	4.5	44
363	Adipose tissue depots of Holstein cows are immune responsive: inflammatory gene expression in vitro. *Domestic Animal Endocrinology*, **2010**, 38, 168-78	2.3	43
362	Expression of metabolic, tissue remodeling, oxidative stress, and inflammatory pathways in mammary tissue during involution in lactating dairy cows. *Bioinformatics and Biology Insights*, **2010**, 4, 85-97	5.3	43
361	Sustained upregulation of stearoyl-CoA desaturase in bovine mammary tissue with contrasting changes in milk fat synthesis and lipogenic gene networks caused by lipid supplements. *Functional and Integrative Genomics*, **2010**, 10, 561-75	3.8	42
360	Ratio of lysine to methionine alters expression of genes involved in milk protein transcription and translation and mTOR phosphorylation in bovine mammary cells. *Physiological Genomics*, **2014**, 46, 268-75	3.6	41
359	Functional adaptations of the transcriptome to mastitis-causing pathogens: the mammary gland and beyond. *Journal of Mammary Gland Biology and Neoplasia*, **2011**, 16, 305-22	2.4	41
358	SCD1 Alters Long-Chain Fatty Acid (LCFA) Composition and Its Expression Is Directly Regulated by SREBP-1 and PPAR in Dairy Goat Mammary Cells. *Journal of Cellular Physiology*, **2017**, 232, 635-649	7	40
357	TRIENNIAL LACTATION SYMPOSIUM: Nutrigenomics in dairy cows: Nutrients, transcription factors, and techniques. *Journal of Animal Science*, **2015**, 93, 5531-53	0.7	40
356	Genes regulating lipid and protein metabolism are highly expressed in mammary gland of lactating dairy goats. *Functional and Integrative Genomics*, **2015**, 15, 309-21	3.8	39
355	Adipose tissue lipogenic gene networks due to lipid feeding and milk fat depression in lactating cows. *Journal of Dairy Science*, **2009**, 92, 4290-300	4	38
354	Inflammation- and lipid metabolism-related gene network expression in visceral and subcutaneous adipose depots of Holstein cows. *Journal of Dairy Science*, **2014**, 97, 3441-8	4	37
353	Dietary lipid during the transition period to manipulate subcutaneous adipose tissue peroxisome proliferator-activated receptor- co-regulator and target gene expression. *Journal of Dairy Science*, **2011**, 94, 5913-25	4	37
352	Methionine and Choline Supply during the Periparturient Period Alter Plasma Amino Acid and One-Carbon Metabolism Profiles to Various Extents: Potential Role in Hepatic Metabolism and Antioxidant Status. *Nutrients*, **2016**, 9,	6.7	36

351	Effects of Dietary l-Arginine and N-Carbamylglutamate Supplementation on Intestinal Integrity, Immune Function, and Oxidative Status in Intrauterine-Growth-Retarded Suckling Lambs. *Journal of Agricultural and Food Chemistry*, **2018**, 66, 4145-4154	5.7	36
350	Gut response induced by weaning in piglet features marked changes in immune and inflammatory response. *Functional and Integrative Genomics*, **2014**, 14, 657-71	3.8	36
349	Hepatic Metabolic, Inflammatory, and Stress-Related Gene Expression in Growing Mice Consuming a Low Dose of Trans-10, cis-12-Conjugated Linoleic Acid. *Journal of Lipids*, **2012**, 2012, 571281	2.7	36
348	High-starch diets induce precocious adipogenic gene network up-regulation in longissimus lumborum of early-weaned Angus cattle. *British Journal of Nutrition*, **2010**, 103, 953-63	3.6	36
347	Glucose transporter and hypoxia-associated gene expression in the mammary gland of transition dairy cattle. *Journal of Dairy Science*, **2011**, 94, 2912-22	4	35
346	Effects of dietary cis 9, trans 11-18:2, trans 10, cis 12-18:2, or vaccenic acid (trans 11-18:1) during lactation on body composition, tissue fatty acid profiles, and litter growth in mice. *British Journal of Nutrition*, **2003**, 90, 1039-48	3.6	35
345	Effects of Arginine concentration on the in vitro expression of Casein and mTOR pathway related genes in mammary epithelial cells from dairy cattle. *PLoS ONE*, **2014**, 9, e95985	3.7	35
344	MicroRNA-26a/b and their host genes synergistically regulate triacylglycerol synthesis by targeting the INSIG1 gene. *RNA Biology*, **2016**, 13, 500-10	4.8	34
343	MicroRNA Bta-miR-181a regulates the biosynthesis of bovine milk fat by targeting ACSL1. *Journal of Dairy Science*, **2016**, 99, 3916-3924	4	34
342	Application of Top-Down and Bottom-up Systems Approaches in Ruminant Physiology and Metabolism. *Current Genomics*, **2012**, 13, 379-94	2.6	34
341	Long-chain fatty acid effects on peroxisome proliferator-activated receptor-alpha-regulated genes in Madin-Darby bovine kidney cells: optimization of culture conditions using palmitate. *Journal of Dairy Science*, **2009**, 92, 2027-37	4	34
340	Grazing allowance after the morning or afternoon milking for lactating cows fed a total mixed ration (TMR) enhances trans11-18:1 and cis9,trans11-18:2 (rumenic acid) in milk fat to different extents. *Animal Feed Science and Technology*, **2003**, 109, 105-119	3	34
339	Dietary impacts on rumen microbiota in beef and dairy production. *Animal Frontiers*, **2016**, 6, 22-29	5.5	34
338	Parturition in dairy cows temporarily alters the expression of genes in circulating neutrophils. *Journal of Dairy Science*, **2016**, 99, 6470-6483	4	34
337	Body condition score and plane of nutrition prepartum affect adipose tissue transcriptome regulators of metabolism and inflammation in grazing dairy cows during the transition period. *Journal of Dairy Science*, **2016**, 99, 758-70	4	33
336	Short communication: Endoplasmic reticulum stress gene network expression in bovine mammary tissue during the lactation cycle. *Journal of Dairy Science*, **2012**, 95, 2562-6	4	33
335	Mammary gene expression profiles during an intramammary challenge reveal potential mechanisms linking negative energy balance with impaired immune response. *Physiological Genomics*, **2010**, 41, 161-70	3.6	33
334	Internal controls for quantitative polymerase chain reaction of swine mammary glands during pregnancy and lactation. *Journal of Dairy Science*, **2008**, 91, 3057-66	4	33

333	Fatty acid-induced endoplasmic reticulum stress promoted lipid accumulation in calf hepatocytes, and endoplasmic reticulum stress existed in the liver of severe fatty liver cows. *Journal of Dairy Science*, **2019**, 102, 7359-7370	4	32
332	Effect of the level of maternal energy intake prepartum on immunometabolic markers, polymorphonuclear leukocyte function, and neutrophil gene network expression in neonatal Holstein heifer calves. *Journal of Dairy Science*, **2013**, 96, 3573-87	4	32
331	Hepatic Activity and Transcription of Betaine-Homocysteine Methyltransferase, Methionine Synthase, and Cystathionine Synthase in Periparturient Dairy Cows Are Altered to Different Extents by Supply of Methionine and Choline. *Journal of Nutrition*, **2017**, 147, 11-19	4.1	32
330	Overfeeding energy upregulates peroxisome proliferator-activated receptor (PPAR)□controlled adipogenic and lipolytic gene networks but does not affect proinflammatory markers in visceral and subcutaneous adipose depots of Holstein cows. *Journal of Dairy Science*, **2014**, 97, 3431-40	4	32
329	Prepartum dietary energy intake alters adipose tissue transcriptome profiles during the periparturient period in Holstein dairy cows. *Journal of Animal Science and Biotechnology*, **2020**, 11, 1	6	32
328	Supplementation with rumen-protected methionine or choline during the transition period influences whole-blood immune response in periparturient dairy cows. *Journal of Dairy Science*, **2017**, 100, 3958-3968	4	31
327	Varying the ratio of Lys:Met while maintaining the ratios of Thr:Phe, Lys:Thr, Lys:His, and Lys:Val alters mammary cellular metabolites, mammalian target of rapamycin signaling, and gene transcription. *Journal of Dairy Science*, **2018**, 101, 1708-1718	4	31
326	Gene expression ratio stability evaluation in prepubertal bovine mammary tissue from calves fed different milk replacers reveals novel internal controls for quantitative polymerase chain reaction. *Journal of Nutrition*, **2008**, 138, 1158-64	4.1	31
325	Integrative analyses of hepatic differentially expressed genes and blood biomarkers during the peripartal period between dairy cows overfed or restricted-fed energy prepartum. *PLoS ONE*, **2014**, 9, e99757	3.7	31
324	Requirement for digestible calcium by eleven- to twenty-five-kilogram pigs as determined by growth performance, bone ash concentration, calcium and phosphorus balances, and expression of genes involved in transport of calcium in intestinal and kidney cells. *Journal of Animal Science*, **2016**, 94, 3321-3334	0.7	31
323	Linking Peripartal Dynamics of Ruminal Microbiota to Dietary Changes and Production Parameters. *Frontiers in Microbiology*, **2016**, 7, 2143	5.7	30
322	Hepatic global DNA and peroxisome proliferator-activated receptor alpha promoter methylation are altered in peripartal dairy cows fed rumen-protected methionine. *Journal of Dairy Science*, **2016**, 99, 234-44	4	29
321	Fertility and the transition dairy cow. *Reproduction, Fertility and Development*, **2017**, 30, 85-100	1.8	29
320	Grazing dairy cows had decreased interferon-□tumor necrosis factor, and interleukin-17, and increased expression of interleukin-10 during the first week after calving. *Journal of Dairy Science*, **2015**, 98, 937-46	4	28
319	Postpartal immunometabolic gene network expression and function in blood neutrophils are altered in response to prepartal energy intake and postpartal intramammary inflammatory challenge. *Journal of Dairy Science*, **2014**, 97, 2165-77	4	28
318	Trans and conjugated fatty acids in milk from cows and goats consuming pasture or receiving vegetable oils or seeds. *Italian Journal of Animal Science*, **2002**, 1, 243-254	2.2	28
317	Feed restriction, but not l-carnitine infusion, alters the liver transcriptome by inhibiting sterol synthesis and mitochondrial oxidative phosphorylation and increasing gluconeogenesis in mid-lactation dairy cows. *Journal of Dairy Science*, **2013**, 96, 2201-2213	4	27
316	Central Role of the PPAR□Gene Network in Coordinating Beef Cattle Intramuscular Adipogenesis in Response to Weaning Age and Nutrition. *Gene Regulation and Systems Biology*, **2014**, 8, 17-32	2	27

315	miR-30e-5p and miR-15a Synergistically Regulate Fatty Acid Metabolism in Goat Mammary Epithelial Cells via LRP6 and YAP1. *International Journal of Molecular Sciences*, **2016**, 17,	6.3	27
314	Fatty acid elongase 6 plays a role in the synthesis of long-chain fatty acids in goat mammary epithelial cells. *Journal of Dairy Science*, **2017**, 100, 4987-4995	4	26
313	Cyanidin-3-O-glucoside improves non-alcoholic fatty liver disease by promoting PINK1-mediated mitophagy in mice. *British Journal of Pharmacology*, **2020**, 177, 3591-3607	8.6	26
312	Essential amino acid ratios and mTOR affect lipogenic gene networks and miRNA expression in bovine mammary epithelial cells. *Journal of Animal Science and Biotechnology*, **2016**, 7, 44	6	26
311	Peroxisome proliferator-activated receptor 1 and 2 isoforms alter lipogenic gene networks in goat mammary epithelial cells to different extents. *Journal of Dairy Science*, **2014**, 97, 5437-47	4	26
310	Short communication: Diurnal profiles of conjugated linoleic acids and trans fatty acids in ruminal fluid from cows fed a high concentrate diet supplemented with fish oil, linseed oil, or sunflower oil. *Journal of Dairy Science*, **2004**, 87, 2468-71	4	26
309	Alterations in blood plasma and milk fatty acid profiles of lactating Holstein cows in response to ruminal infusion of a conjugated linoleic acid mixture. *Animal Research*, **2001**, 50, 463-476		26
308	MicroRNA-106b Regulates Milk Fat Metabolism via ATP Binding Cassette Subfamily A Member 1 (ABCA1) in Bovine Mammary Epithelial Cells. *Journal of Agricultural and Food Chemistry*, **2019**, 67, 3981-3990	5.7	25
307	Adipose and liver gene expression profiles in response to treatment with a nonsteroidal antiinflammatory drug after calving in grazing dairy cows. *Journal of Dairy Science*, **2015**, 98, 3079-85	4	25
306	Overfeeding Dairy Cattle During Late-Pregnancy Alters Hepatic PPAR Regulated Pathways Including Hepatokines: Impact on Metabolism and Peripheral Insulin Sensitivity. *Gene Regulation and Systems Biology*, **2014**, 8, 97-111	2	25
305	Ruminant metabolic systems biology: reconstruction and integration of transcriptome dynamics underlying functional responses of tissues to nutrition and physiological state. *Gene Regulation and Systems Biology*, **2012**, 6, 109-25	2	25
304	Dietary trans-vaccenic acid (trans11-18:1) increases concentration of cis9,transll-conjugated linoleic acid (rumenic acid) in tissues of lactating mice and suckling pups. *Reproduction, Nutrition, Development*, **2002**, 42, 85-99		25
303	Enhanced supply of methionine or arginine alters mechanistic target of rapamycin signaling proteins, messenger RNA, and microRNA abundance in heat-stressed bovine mammary epithelial cells in vitro. *Journal of Dairy Science*, **2019**, 102, 2469-2480	4	25
302	Prepartal standing behavior as a parameter for early detection of postpartal subclinical ketosis associated with inflammation and liver function biomarkers in peripartal dairy cows. *Journal of Dairy Science*, **2018**, 101, 8224-8235	4	25
301	Prepartal dietary energy level affects peripartal bovine blood neutrophil metabolic, antioxidant, and inflammatory gene expression. *Journal of Dairy Science*, **2015**, 98, 5492-505	4	24
300	Effect of dietary starch level and high rumen-undegradable protein on endocrine-metabolic status, milk yield, and milk composition in dairy cows during early and late lactation. *Journal of Dairy Science*, **2014**, 97, 7788-803	4	24
299	Effects of the peroxisome proliferator-activated receptor-alpha agonists clofibrate and fish oil on hepatic fatty acid metabolism in weaned dairy calves. *Journal of Dairy Science*, **2010**, 93, 2404-18	4	24
298	Functional and gene network analyses of transcriptional signatures characterizing pre-weaned bovine mammary parenchyma or fat pad uncovered novel inter-tissue signaling networks during development. *BMC Genomics*, **2010**, 11, 331	4.5	24

297	Tea polyphenols protect bovine mammary epithelial cells from hydrogen peroxide-induced oxidative damage in vitro by activating NFE2L2/HMOX1 pathways. *Journal of Dairy Science*, **2019**, 102, 1658-1670	4	24
296	Methionine and valine activate the mammalian target of rapamycin complex 1 pathway through heterodimeric amino acid taste receptor (TAS1R1/TAS1R3) and intracellular Ca in bovine mammary epithelial cells. *Journal of Dairy Science*, **2018**, 101, 11354-11363	4	24
295	Maternal supplementation with rumen-protected methionine increases prepartal plasma methionine concentration and alters hepatic mRNA abundance of 1-carbon, methionine, and transsulfuration pathways in neonatal Holstein calves. *Journal of Dairy Science*, **2017**, 100, 3209-3219	4	23
294	Supplemental methionine, choline, or taurine alter in vitro gene network expression of polymorphonuclear leukocytes from neonatal Holstein calves. *Journal of Dairy Science*, **2017**, 100, 3155-3165	4	23
293	Innate immune responses induced by lipopolysaccharide and lipoteichoic acid in primary goat mammary epithelial cells. *Journal of Animal Science and Biotechnology*, **2017**, 8, 29	6	23
292	TRIENNIAL LACTATION SYMPOSIUM: Nutrigenomics in livestock: Systems biology meets nutrition. *Journal of Animal Science*, **2015**, 93, 5554-74	0.7	23
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