

Penny K Riggs

List of Publications by Year in Descending Order

Source: <https://exaly.com/author-pdf/8758673/penny-k-riggs-publications-by-year.pdf>

Version: 2024-04-28

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

78
papers

2,287
citations

20
h-index

47
g-index

85
ext. papers

2,745
ext. citations

3.1
avg. IF

4.74
L-index

#	Paper	IF	Citations
78	Skeletal Muscle Expression of Actinin-3 () in Relation to Feed Efficiency Phenotype of F - Steers.. <i>Frontiers in Genetics</i> , 2022 , 13, 796038	4.5	1
77	Differential Expression of MicroRNAs in Dark-Cutting Meat from Beef Carcasses. <i>Applied Sciences (Switzerland)</i> , 2022 , 12, 3555	2.6	0
76	The Potential Use of DNA Methylation Patterns from Peripheral Blood Leukocytes as a Surrogate for Stress Axis Tissues in Mature Brahman Cows. <i>Journal of Animal Science</i> , 2021 , 99, 1-2	0.7	
75	Evaluation of Prenatal Transportation Stress on the Number of Pituitary Corticotrophs in Mature Brahman Cows. <i>Journal of Animal Science</i> , 2021 , 99, 24-25	0.7	78
74	Evaluation of Peripheral Blood Leukocytes as a Surrogate for RNA-Seq Detection of Gene Expression in Stress Axis Tissues of Mature Brahman Cows. <i>Journal of Animal Science</i> , 2021 , 99, 2-3	0.7	78
73	Differential Expression of Circadian Clock Genes in the Bovine Neuroendocrine Adrenal System. <i>Journal of the Endocrine Society</i> , 2021 , 5, A66-A67	0.4	
72	Genome-wide DNA methylation alteration in prenatally stressed Brahman heifer calves with the advancement of age. <i>Epigenetics</i> , 2021 , 16, 519-536	5.7	2
71	Distributional characterizations and testing for differences of relatedness and inbreeding of a subpopulation of American Hereford bulls. <i>Translational Animal Science</i> , 2021 , 5, txab008	1.4	
70	Genome association of carcass and palatability traits from Bos indicus-Bos taurus crossbred steers within electrical stimulation status and correspondence with steer temperament 3. Aroma and flavor attributes of cooked steaks. <i>Livestock Science</i> , 2020 , 233, 103943	1.7	1
69	Genome association of carcass and palatability traits from Bos indicus-Bos taurus crossbred steers within electrical stimulation status and correspondence with steer temperament 2. Palatability. <i>Livestock Science</i> , 2020 , 232, 103897	1.7	
68	Influence of prenatal transportation stress-induced differential DNA methylation on the physiological control of behavior and stress response in suckling Brahman bull calves. <i>Journal of Animal Science</i> , 2020 , 98,	0.7	8
67	Effect of prenatal transportation stress on DNA methylation in Brahman heifers. <i>Livestock Science</i> , 2020 , 240, 104116	1.7	3
66	Genome to Phenome: Improving Animal Health, Production, and Well-Being - A New USDA Blueprint for Animal Genome Research 2018-2027. <i>Frontiers in Genetics</i> , 2019 , 10, 327	4.5	47
65	Expansion of Knowledge and Advances in Genetics for Quantitative Analyses 2019 , 16-27		
64	Considerations for the use of Cre recombinase for conditional gene deletion in the mouse lens. <i>Human Genomics</i> , 2019 , 13, 10	6.8	14
63	376 Awardee Talk - Prenatal transportation stress alters physiology of suckling Brahman calves as mediated by changes in DNA methylation. <i>Journal of Animal Science</i> , 2019 , 97, 142-142	0.7	78
62	PSVIII-40 Effect of prenatal transportation stress on DNA methylation in Brahman heifers. <i>Journal of Animal Science</i> , 2019 , 97, 262-263	0.7	78

61	PSXIII-13 Comparison of telomere length in age-matched primiparous and multiparous Brahman cows. <i>Journal of Animal Science</i> , 2019 , 97, 363-364	0.7	78
60	PSVIII-39 Genome-wide DNA methylation alteration in prenatally stressed Brahman heifer calves with the advancement of age. <i>Journal of Animal Science</i> , 2019 , 97, 263-264	0.7	78
59	Genome association of carcass and palatability traits from Bos indicus-Bos taurus crossbred steers within electrical stimulation status and correspondence with steer temperament 1. Carcass. <i>Livestock Science</i> , 2019 , 229, 150-158	1.7	4
58	Prenatal transportation stress alters genome-wide DNA methylation in suckling Brahman bull calves. <i>Journal of Animal Science</i> , 2018 , 96, 5075-5099	0.7	17
57	The outbreak that changed meat and poultry inspection systems worldwide. <i>Animal Frontiers</i> , 2018 , 8, 4-8	5.5	3
56	Food and nutrient security for a growing population. <i>Animal Frontiers</i> , 2018 , 8, 3-4	5.5	4
55	Association of udder traits with single nucleotide polymorphisms in crossbred - cows. <i>Journal of Animal Science</i> , 2017 , 95, 2399-2407	0.7	9
54	Association of udder traits with single nucleotide polymorphisms in crossbred cows. <i>Journal of Animal Science</i> , 2017 , 95, 2399	0.7	7
53	Crossbred steer temperament as yearlings and whole genome association of steer temperament as yearlings and calf temperament post-weaning. <i>Journal of Animal Science</i> , 2016 , 94, 1408-14	0.7	10
52	003 Genome wide association of beef flavor and tenderness in steaks (with electrical stimulation and without) from carcasses of Nellore-Bos taurus steers. <i>Journal of Animal Science</i> , 2016 , 94, 2-2	0.7	1
51	095 Metabolomic profiling for identification of biomarkers associated with temperament in feedlot cattle. <i>Journal of Animal Science</i> , 2016 , 94, 47-47	0.7	
50	Development of a 63K SNP Array for Cotton and High-Density Mapping of Intraspecific and Interspecific Populations of Gossypium spp. <i>G3: Genes, Genomes, Genetics</i> , 2015 , 5, 1187-209	3.2	117
49	Constitutive activation of Stat3 in mouse epidermis is linked to hair deficiency and cytoskeletal network damage. <i>Experimental Dermatology</i> , 2015 , 24, 796-8	4	1
48	Differential miRNA expression in inherently high- and low-active inbred mice. <i>Physiological Reports</i> , 2015 , 3, e12469	2.6	12
47	Description of bovine major histocompatibility complex class IIa haplotypes using parthenogenetic embryo-derived cells. <i>Animal Genetics</i> , 2015 , 46, 325-8	2.5	
46	Dexamethasone acutely down-regulates genes involved in steroidogenesis in stallion testes. <i>Journal of Steroid Biochemistry and Molecular Biology</i> , 2014 , 143, 451-9	5.1	23
45	Expression of bovine genes associated with local and systemic immune response to infestation with the Lone Star tick, <i>Amblyomma americanum</i> . <i>Ticks and Tick-borne Diseases</i> , 2014 , 5, 676-88	3.6	3
44	Genetic evaluation of aspects of temperament in Nellore-Angus calves. <i>Journal of Animal Science</i> , 2014 , 92, 3223-30	0.7	10

43	Differential gene expression in high- and low-active inbred mice. <i>BioMed Research International</i> , 2014 , 2014, 361048	3	8
42	Alternative parameterizations of relatedness in whole genome association analysis of pre-weaning traits of Nelore-Angus calves. <i>Genetics and Molecular Biology</i> , 2014 , 37, 518-25	2	3
41	Marinobufagenin regulates permeability and gene expression of brain endothelial cells. <i>American Journal of Physiology - Regulatory Integrative and Comparative Physiology</i> , 2014 , 306, R918-24	3.2	14
40	Consequences of perinatal bisphenol A exposure in a mouse model of multiple sclerosis. <i>Autoimmunity</i> , 2014 , 47, 57-66	3	15
39	Fine mapping reveals that promotion susceptibility locus 1 (Psl1) is a compound locus with multiple genes that modify susceptibility to skin tumor development. <i>G3: Genes, Genomes, Genetics</i> , 2014 , 4, 1071-7	3.3	2
38	Hypothalamic distribution, adenohipophyseal receptor expression, and ligand functionality of RFamide-related peptide 3 in the mare during the breeding and nonbreeding seasons. <i>Biology of Reproduction</i> , 2014 , 90, 28	3.9	26
37	Genome-wide association study of temperament and tenderness using different Bayesian approaches in a Nellore-Angus crossbred population. <i>Livestock Science</i> , 2014 , 161, 17-27	1.7	19
36	A high-resolution radiation hybrid map of the river buffalo major histocompatibility complex and comparison with BoLA. <i>Animal Genetics</i> , 2013 , 44, 369-76	2.5	2
35	Evaluation of methods for the isolation of high quality RNA from bovine and cervine hide biopsies. <i>Journal of Parasitology</i> , 2013 , 99, 19-23	0.9	1
34	Development and characterization of two porcine monocyte-derived macrophage cell lines. <i>Results in Immunology</i> , 2013 , 3, 26-32		12
33	Whole genome association of SNP with newborn calf cannon bone length. <i>Livestock Science</i> , 2013 , 155, 186-196	1.7	7
32	Gene expression in the arcuate nucleus of heifers is affected by controlled intake of high- and low-concentrate diets. <i>Journal of Animal Science</i> , 2012 , 90, 2222-32	0.7	27
31	Proteomic and pathway analyses reveal a network of inflammatory genes associated with differences in skin tumor promotion susceptibility in DBA/2 and C57BL/6 mice. <i>Carcinogenesis</i> , 2012 , 33, 2208-19	4.6	15
30	Expression of Hypothalamic RF-amide Related Peptide 3 (RFRP3) and Adenohipophyseal G Protein-Coupled Receptor 147 (GPR147) During the Breeding and Non-breeding Seasons in the Mare.. <i>Biology of Reproduction</i> , 2012 , 87, 482-482	3.9	
29	Evidence that Gsta4 modifies susceptibility to skin tumor development in mice and humans. <i>Journal of the National Cancer Institute</i> , 2010 , 102, 1663-75	9.7	18
28	Discovery of candidate genes and pathways in the endometrium regulating ovine blastocyst growth and conceptus elongation. <i>Physiological Genomics</i> , 2009 , 39, 85-99	3.6	66
27	Fragile sites in domestic animal chromosomes: molecular insights and challenges. <i>Cytogenetic and Genome Research</i> , 2009 , 126, 97-109	1.9	8
26	The genome sequence of taurine cattle: a window to ruminant biology and evolution. <i>Science</i> , 2009 , 324, 522-8	33.3	863

25	First radiation hybrid map of the river buffalo X chromosome (BBUX) and comparison with BTAX. <i>Animal Genetics</i> , 2008 , 39, 196-200	2.5	8
24	Application of dissociation curve analysis to radiation hybrid panel marker scoring: generation of a map of river buffalo (<i>B. bubalis</i>) chromosome 20. <i>BMC Genomics</i> , 2008 , 9, 544	4.5	2
23	A first generation whole genome RH map of the river buffalo with comparison to domestic cattle. <i>BMC Genomics</i> , 2008 , 9, 631	4.5	50
22	Progressive metaplastic and dysplastic changes in mouse pancreas induced by cyclooxygenase-2 overexpression. <i>Neoplasia</i> , 2008 , 10, 782-96	6.4	69
21	Report of a chimeric origin of transposable elements in a bovine-coding gene. <i>Genetics and Molecular Research</i> , 2008 , 7, 107-16	1.2	1
20	Mapping MHC genes in river buffalo. <i>Developments in Biologicals</i> , 2008 , 132, 343-346		4
19	Identification of Progesterone-Regulated Genes Governing Pre-implantation Conceptus Growth and Development.. <i>Biology of Reproduction</i> , 2008 , 78, 173-173	3.9	1
18	Differential expression of multiple anti-apoptotic proteins in epidermis of IGF-1 transgenic mice as revealed by 2-dimensional gel electrophoresis/mass spectrometry analysis. <i>Molecular Carcinogenesis</i> , 2007 , 46, 331-40	5	10
17	Preliminary radiation hybrid map for river buffalo chromosome 6 and comparison to bovine chromosome 3. <i>Animal Genetics</i> , 2007 , 38, 406-9	2.5	11
16	The contribution of transposable elements to <i>Bos taurus</i> gene structure. <i>Gene</i> , 2007 , 390, 180-9	3.8	24
15	A radiation hybrid map of river buffalo (<i>Bubalus bubalis</i>) chromosome 1 (BBU1). <i>Cytogenetic and Genome Research</i> , 2007 , 119, 100-4	1.9	13
14	Ionizing radiation-induced bioeffects in space and strategies to reduce cellular injury and carcinogenesis. <i>Aviation, Space, and Environmental Medicine</i> , 2007 , 78, A67-78		8
13	ATM controls c-Myc and DNA synthesis during postnatal thymocyte development through regulation of redox state. <i>Free Radical Biology and Medicine</i> , 2006 , 41, 640-8	7.8	15
12	Differential gene expression in epidermis of mice sensitive and resistant to phorbol ester skin tumor promotion. <i>Molecular Carcinogenesis</i> , 2005 , 44, 122-36	5	28
11	Alteration of Egr-1 mRNA during multistage carcinogenesis in mouse skin. <i>Molecular Carcinogenesis</i> , 2000 , 27, 247-51	5	22
10	Preparation of Metaphase Chromosomes for Cytogenetic Analysis 1998 , 229-238		
9	Development and initial characterization of a <i>Bos taurus</i> x <i>B. gaurus</i> interspecific hybrid backcross panel. <i>Journal of Heredity</i> , 1997 , 88, 373-9	2.4	21
8	A high resolution GBG-banded karyotype of the Atlantic bottlenose dolphin, <i>Tursiops truncatus</i> : generation of an ideogram, and NOR localization by fluorescence in situ hybridization. <i>Cytogenetic and Genome Research</i> , 1997 , 78, 6-11	1.9	6

7	Structure of the human gene (COX6A2) for the heart/muscle isoform of cytochrome c oxidase subunit VIa and its chromosomal location in humans, mice, and cattle. <i>Genomics</i> , 1997 , 42, 146-51	4.3	11
6	Standardization of cattle karyotype nomenclature: report of the committee for the standardization of the cattle karyotype. <i>Cytogenetic and Genome Research</i> , 1996 , 74, 259-61	1.9	67
5	Chromosomal localization of the porcine skeletal muscle calpain gene. <i>Mammalian Genome</i> , 1996 , 7, 226-8	3.2	12
4	Structure and chromosomal location of the bovine gene for the heart muscle isoform of cytochrome c oxidase subunit VIII. <i>Mammalian Genome</i> , 1995 , 6, 118-22	3.2	11
3	Analysis of aphidicolin-induced chromosome fragility in the domestic pig (<i>Sus scrofa</i>). <i>Cytogenetic and Genome Research</i> , 1993 , 62, 110-6	1.9	19
2	Identification of aphidicolin-induced fragile sites in domestic pig chromosomes. <i>Genetics Selection Evolution</i> , 1991 , 23, S187	4.9	3
1	Chromosomes of Fish. <i>Advances in Veterinary Medicine</i> , 1990 , 34, 209-227		