Kieran G Meade

List of Publications by Year in descending order

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172457 197818 2,716 82 29 49 citations h-index g-index papers 85 85 85 2876 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Induction of a Novel Chicken Toll-Like Receptor following <i>Salmonella enterica</i> Serovar Typhimurium Infection. Infection and Immunity, 2006, 74, 1692-1698.	2.2	173
2	Toll-like receptor and antimicrobial peptide expression in the bovine endometrium. Reproductive Biology and Endocrinology, 2008, 6, 53.	3.3	167
3	Histopathological and molecular evaluation of Holstein-Friesian cows postpartum: Toward an improved understanding of uterine innate immunity. Theriogenology, 2009, 71, 1396-1407.	2.1	132
4	\hat{l}^2 -Defensins: Farming the Microbiome for Homeostasis and Health. Frontiers in Immunology, 2018, 9, 3072.	4.8	111
5	Tuberculosis Immunity: Opportunities from Studies with Cattle. Clinical and Developmental Immunology, 2011, 2011, 1-11.	3.3	104
6	Comparative in vivo infection models yield insights on early host immune response to Campylobacter in chickens. Immunogenetics, 2009, 61, 101-110.	2.4	92
7	Experimental Staphylococcus aureus infection of the mammary gland induces region-specific changes in innate immune gene expression. Veterinary Immunology and Immunopathology, 2011, 140, 181-189.	1.2	87
8	Bovine \hat{l}^2 -defensin gene family: opportunities to improve animal health? Physiological Genomics, 2014, 46, 17-28.	2.3	81
9	Tumour necrosis factor-α (TNF-α) increases nuclear factor ΰB (NFΰB) activity in and interleukin-8 (IL-8) release from bovine mammary epithelial cells. Veterinary Immunology and Immunopathology, 2007, 116, 59-68.	1.2	77
10	Evolution, expression and effectiveness in a cluster of novel bovine \hat{l}^2 -defensins. Immunogenetics, 2008, 60, 147-156.	2.4	73
11	Innate immune gene expression differentiates the early avian intestinal response between Salmonella and Campylobacter. Veterinary Immunology and Immunopathology, 2009, 132, 191-198.	1.2	71
12	Innate gene repression associated with Mycobacterium bovis infection in cattle: toward a gene signature of disease. BMC Genomics, 2007, 8, 400.	2.8	65
13	RNA-seq Transcriptional Profiling of Peripheral Blood Leukocytes from Cattle Infected with Mycobacterium bovis. Frontiers in Immunology, 2014, 5, 396.	4.8	65
14	Differential antimicrobial peptide gene expression patterns during early chicken embryological development. Developmental and Comparative Immunology, 2009, 33, 516-524.	2.3	64
15	The postpartum endometrial inflammatory response: a normal physiological event with potential implications for bovine fertility. Reproduction, Fertility and Development, 2012, 24, 1028.	0.4	62
16	Analysis of the Bovine Monocyte-Derived Macrophage Response to Mycobacterium avium Subspecies Paratuberculosis Infection Using RNA-seq. Frontiers in Immunology, 2015, 6, 23.	4.8	61
17	Endometrial biopsy: a valuable clinical and research tool in bovine reproduction. Theriogenology, 2010, 73, 988-994.	2.1	57
18	Single Nucleotide Polymorphisms in the Insulin-Like Growth Factor 1 (IGF-1) Gene are Associated with Performance in Holstein-Friesian Dairy Cattle. Frontiers in Genetics, 2011, 2, 3.	2.3	50

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19	Endometrial epithelial cells are potent producers of tracheal antimicrobial peptide and serum amyloid A3 gene expression in response to E. coli stimulation. Veterinary Immunology and Immunopathology, 2013, 151, 157-162.	1.2	50
20	Cytokine mRNA profiling of peripheral blood mononuclear cells from trypanotolerant and trypanosusceptible cattle infected with Trypanosoma congolense. Physiological Genomics, 2006, 28, 53-61.	2.3	49
21	Avian Resistance to Campylobacter jejuni Colonization Is Associated with an Intestinal Immunogene Expression Signature Identified by mRNA Sequencing. PLoS ONE, 2012, 7, e40409.	2.5	46
22	Cauda Epididymis-Specific Beta-Defensin 126 Promotes Sperm Motility but Not Fertilizing Ability in Cattle. Biology of Reproduction, 2016, 95, 122-122.	2.7	44
23	Transcriptional profiling of cattle infected with Trypanosoma congolense highlights gene expression signatures underlying trypanotolerance and trypanosusceptibility. BMC Genomics, 2009, 10, 207.	2.8	41
24	Genome-wide transcriptional profiling of peripheral blood leukocytes from cattle infected with Mycobacterium bovis reveals suppression of host immune genes. BMC Genomics, 2011, 12, 611.	2.8	40
25	Non-canonical Inflammasome-Mediated IL- $1\hat{l}^2$ Production by Primary Endometrial Epithelial and Stromal Fibroblast Cells Is NLRP3 and Caspase-4 Dependent. Frontiers in Immunology, 2019, 10, 102.	4.8	37
26	Reproductive tissue-specific expression profiling and genetic variation across a 19 gene bovine \hat{l}^2 -defensin cluster. Immunogenetics, 2011, 63, 641-651.	2.4	33
27	Characterisation and expression profile of the bovine cathelicidin gene repertoire in mammary tissue. BMC Genomics, 2014, 15, 128.	2.8	33
28	Integrated analysis of the local and systemic changes preceding the development of post-partum cytological endometritis. BMC Genomics, 2015, 16, 811.	2.8	33
29	Cervico-vaginal mucus (CVM) – an accessible source of immunologically informative biomolecules. Veterinary Research Communications, 2018, 42, 255-263.	1.6	33
30	Epigenetic regulation of the innate immune response to LPS in bovine peripheral blood mononuclear cells (PBMC). Veterinary Immunology and Immunopathology, 2013, 154, 102-110.	1,2	32
31	Gene expression profiling of peripheral blood mononuclear cells (PBMC) from Mycobacterium bovis infected cattle after in vitro antigenic stimulation with purified protein derivative of tuberculin (PPD). Veterinary Immunology and Immunopathology, 2006, 113 , 73 -89.	1.2	30
32	Profiling inflammatory biomarkers in cervico-vaginal mucus (CVM) postpartum: Potential early indicators of bovine clinical endometritis?. Theriogenology, 2017, 103, 117-122.	2.1	30
33	The CD4+ T cell methylome contributes to a distinct CD4+ T cell transcriptional signature in Mycobacterium bovis-infected cattle. Scientific Reports, 2016, 6, 31014.	3.3	28
34	Alum Activates the Bovine NLRP3 Inflammasome. Frontiers in Immunology, 2017, 8, 1494.	4.8	27
35	Short communication: Uncovering quantitative trait loci associated with resistance to Mycobacterium avium ssp. paratuberculosis infection in Holstein cattle using a high-density single nucleotide polymorphism panel. Journal of Dairy Science, 2018, 101, 7280-7286.	3.4	27
36	Global endometrial transcriptomic profiling: transient immune activation precedes tissue proliferation and repair in healthy beef cows. BMC Genomics, 2012, 13, 489.	2.8	26

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37	Sperm-Coating Beta-Defensin 126 Is a Dissociation-Resistant Dimer Produced by Epididymal Epithelium in the Bovine Reproductive Tract. Biology of Reproduction, 2016, 95, 121-121.	2.7	22
38	The genetic architecture of milk ELISA scores as an indicator of Johne's disease (paratuberculosis) in dairy cattle. Journal of Dairy Science, 2018, 101, 10062-10075.	3.4	22
39	Technical note: Comparative analyses of the quality and yield of genomic DNA from invasive and noninvasive, automated and manual extraction methods. Journal of Dairy Science, 2011, 94, 3159-3165.	3.4	21
40	A novel subclass of bovine \hat{l}^2 -defensins links reproduction and immunology. Reproduction, Fertility and Development, 2014, 26, 769.	0.4	21
41	Recombinant \hat{l}^2 -defensin 126 promotes bull sperm binding to bovine oviductal epithelia. Reproduction, Fertility and Development, 2018, 30, 1472.	0.4	21
42	Antigen stimulation of peripheral blood mononuclear cells from Mycobacterium bovis infected cattle yields evidence for a novel gene expression program. BMC Genomics, 2008, 9, 447.	2.8	20
43	Genome-Wide Association Analysis of Avian Resistance to <i>Campylobacter jejuni</i> Colonization Identifies Risk Locus Spanning the <i>CDH13</i> Gene. G3: Genes, Genomes, Genetics, 2013, 3, 881-890.	1.8	20
44	Genomic identification, expression profiling, and functional characterization of CatSper channels in the bovineâ€. Biology of Reproduction, 2017, 97, 302-312.	2.7	20
45	A dual targeted \hat{I}^2 -defensin and exome sequencing approach to identify, validate and functionally characterise genes associated with bull fertility. Scientific Reports, 2017, 7, 12287.	3.3	19
46	Global gene expression analysis of chicken caecal response to Campylobacter jejuni. Veterinary Immunology and Immunopathology, 2011, 142, 64-71.	1.2	18
47	Comparative epigenetics: relevance to the regulation of production and health traits in cattle. Animal Genetics, 2014, 45, 3-14.	1.7	17
48	Functional characterisation of bovine interleukin 8 promoter haplotypes in vitro. Molecular Immunology, 2012, 50, 108-116.	2.2	15
49	Comparative genomic identification and validation of \hat{l}^2 -defensin genes in the Ovis aries genome. BMC Genomics, 2017, 18, 278.	2.8	14
50	Prepartum concentrate supplementation of a diet based on medium-quality grass silage: Effects on performance, health, fertility, metabolic function, and immune function of low body condition score cows. Journal of Dairy Science, 2016, 99, 7102-7122.	3.4	13
51	Johne's Disease in Dairy Cattle: An Immunogenetic Perspective. Frontiers in Veterinary Science, 2021, 8, 718987.	2.2	13
52	Characterization of circulating plasma proteins in dairy cows with cytological endometritis. Journal of Proteomics, 2019, 205, 103421.	2.4	12
53	Divergent antimicrobial peptide (AMP) and acute phase protein (APP) responses to Trypanosoma congolense infection in trypanotolerant and trypanosusceptible cattle. Molecular Immunology, 2009, 47, 196-204.	2.2	11
54	In vivo relevance of polymorphic Interleukin 8 promoter haplotype for the systemic immune response to LPS in Holstein-Friesian calves. Veterinary Immunology and Immunopathology, 2016, 182, 1-10.	1.2	11

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55	Functional analysis of bovine interleukin-10 receptor alpha in response to Mycobacterium avium subsp. paratuberculosis lysate using CRISPR/Cas9. BMC Genetics, 2020, 21, 121.	2.7	11
56	Directed alteration of a novel bovine \hat{l}^2 -defensin to improve antimicrobial efficacy against methicillin-resistant Staphylococcus aureus (MRSA). International Journal of Antimicrobial Agents, 2008, 32, 392-397.	2.5	10
57	The role of oct-1 in the regulation of tracheal antimicrobial peptide (TAP) and lingual antimicrobial peptide (LAP) expression in bovine mammary epithelial cells. Immunogenetics, 2011, 63, 715-725.	2.4	10
58	Comparative genomic identification and expression profiling of a novel \hat{l}^2 -defensin gene cluster in the equine reproductive tract. Reproduction, Fertility and Development, 2016, 28, 1499.	0.4	10
59	Improved detection of biomarkers in cervico-vaginal mucus (CVM) from postpartum cattle. BMC Veterinary Research, 2018, 14, 297.	1.9	10
60	Qualitative and quantitative differences in endometrial inflammatory gene expression precede the development of bovine uterine disease. Scientific Reports, 2020, 10, 18275.	3.3	10
61	Improved filtration method to isolate pure populations of primary bovine endometrial epithelial and stromal cells for immunological studies. Veterinary Research Communications, 2020, 44, 29-39.	1.6	10
62	Characterization of the bovine salivary gland transcriptome associated with Mycobacterium avium subsp. paratuberculosis experimental challenge. BMC Genomics, 2019, 20, 491.	2.8	9
63	Interleukin 8 haplotypes drive divergent responses in uterine endometrial cells and are associated with somatic cell score in Holstein-Friesian cattle. Veterinary Immunology and Immunopathology, 2017, 184, 18-28.	1.2	8
64	Purulent vaginal discharge diagnosed in pasture-based Holstein-Friesian cows at 21 days postpartum is influenced by previous lactation milk yield and results in diminished fertility. Journal of Dairy Science, 2020, 103, 666-675.	3.4	8
65	Low serum vitamin D concentrations in Spring-born dairy calves are associated with elevated peripheral leukocytes. Scientific Reports, 2021, 11, 18969.	3.3	8
66	Conserved and breed-specific differences in the cervical transcriptome of sheep with divergent fertility at the follicular phase of a natural oestrus cycle. BMC Genomics, 2021, 22, 752.	2.8	8
67	Transmission ratio distortion at the growth hormone gene (<i>GH1</i>) in bovine preimplantation embryos: An in vitro cultureâ€induced phenomenon?. Molecular Reproduction and Development, 2008, 75, 715-722.	2.0	7
68	Integrated analyses of the microbiological, immunological and ontological transitions in the calf ileum during early life. Scientific Reports, 2020, 10, 21264.	3.3	6
69	1,25(OH)D vitamin D promotes NOS2 expression in response to bacterial and viral PAMPs in primary bovine salivary gland fibroblasts. Veterinary Research Communications, 2020, 44, 83-88.	1.6	6
70	Biochemical and molecular characterization of sialylated cervical mucins in sheep. Biology of Reproduction, 2022, 107, 419-431.	2.7	6
71	Application of the TruCulture \hat{A}^{\odot} whole blood stimulation system for immune response profiling in cattle. Veterinary Immunology and Immunopathology, 2020, 221, 110025.	1.2	5
72	Bovine innate immune phenotyping via a standardized whole blood stimulation assay. Scientific Reports, 2021, 11, 17227.	3.3	5

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73	Effect of IL-8 haplotype on temporal profile in circulating concentrations of interleukin 8 and 25(OH) vitamin D in Holstein-Friesian calves. Veterinary Immunology and Immunopathology, 2021, 238, 110287.	1.2	4
74	Ewe breed differences in the cervical transcriptome at the follicular phase of a synchronised oestrous cycle. BMC Genomics, 2022, 23, 363.	2.8	4
75	The immune response in bovine primary dermal fibroblasts is influenced by Interleukin 8 promoter haplotype and vitamin D. Veterinary Immunology and Immunopathology, 2021, 238, 110291.	1.2	3
76	Advances in Bovine Immunology ââ,¬â€œ New Tools and New Insights to Tackle Old Foes. Frontiers in Immunology, 2015, 6, 71.	4.8	2
77	Association of genetic polymorphisms related to Johne's disease with estimated breeding values of Holstein sires for milk ELISA test scores. BMC Veterinary Research, 2020, 16, 165.	1.9	2
78	Effect of IL8 haplotype on immunological traits in periparturient dairy cows. Veterinary Immunology and Immunopathology, 2021, 238, 110288.	1,2	2
79	A preliminary analysis of the variation in circulating 25-hydroxycholecalciferol concentrations in peri-partum spring-calving dairy cows. Veterinary Research Communications, 0, , .	1.6	1
80	Convenient detection of single nucleotide polymorphism haplotypes in the bovine growth hormone gene using amplification-created restriction sites. Animal Genetics, 2005, 36, 175-177.	1.7	0
81	3 ROLE OF \hat{I}^2 -DEFENSIN 126 IN PROMOTING SPERM MOTILITY IN CATTLE. Reproduction, Fertility and Development, 2016, 28, 131.	0.4	0
82	Cervical immune activation during the luteal phase may compromise subsequent trans-cervical ram sperm transport. Biology of Reproduction, 0, , .	2.7	0