

Elizabeth J Glass

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5354903/publications.pdf>

Version: 2024-02-01

86
papers

4,504
citations

101496

36
h-index

106281

65
g-index

87
all docs

87
docs citations

87
times ranked

5293
citing authors

#	ARTICLE	IF	CITATIONS
1	Enhancing the toolbox to study IL-17A in cattle and sheep. <i>Veterinary Research</i> , 2017, 48, 20.	1.1	17
2	Host species adaptation of TLR5 signalling and flagellin recognition. <i>Scientific Reports</i> , 2017, 7, 17677.	1.6	27
3	Live and inactivated <i>Salmonella enterica</i> serovar Typhimurium stimulate similar but distinct transcriptome profiles in bovine macrophages and dendritic cells. <i>Veterinary Research</i> , 2016, 47, 46.	1.1	9
4	Phenotypic and functional analysis of monocyte populations in cattle peripheral blood identifies a subset with high endocytic and allogeneic T-cell stimulatory capacity. <i>Veterinary Research</i> , 2015, 46, 112.	1.1	49
5	Transcriptomic Profiling of Virus-Host Cell Interactions following Chicken Anaemia Virus (CAV) Infection in an In Vivo Model. <i>PLoS ONE</i> , 2015, 10, e0134866.	1.1	19
6	Functional analysis of bovine TLR5 and association with IgA responses of cattle following systemic immunisation with H7 flagella. <i>Veterinary Research</i> , 2015, 46, 9.	1.1	17
7	Late production of CXCL8 in ruminant oro-nasal turbinate cells in response to <i>Chlamydia abortus</i> infection. <i>Veterinary Immunology and Immunopathology</i> , 2015, 168, 97-102.	0.5	5
8	Genomic Prediction for Tuberculosis Resistance in Dairy Cattle. <i>PLoS ONE</i> , 2014, 9, e96728.	1.1	42
9	Genome-wide association study identifies novel loci associated with resistance to bovine tuberculosis. <i>Heredity</i> , 2014, 112, 543-551.	1.2	92
10	The level of H ₂ O ₂ type oxidative stress regulates virulence of <i>Theileria</i> transformed leukocytes. <i>Cellular Microbiology</i> , 2014, 16, 269-279.	1.1	34
11	Differential response of bovine mammary epithelial cells to <i>Staphylococcus aureus</i> or <i>Escherichia coli</i> agonists of the innate immune system. <i>Veterinary Research</i> , 2013, 44, 40.	1.1	191
12	Detectability of bovine TB using the tuberculin skin test does not vary significantly according to pathogen genotype within Northern Ireland. <i>Infection, Genetics and Evolution</i> , 2013, 19, 15-22.	1.0	13
13	<i>Escherichia coli</i> - and <i>Staphylococcus aureus</i> -induced mastitis differentially modulate transcriptional responses in neighbouring uninfected bovine mammary gland quarters. <i>BMC Genomics</i> , 2013, 14, 36.	1.2	125
14	Field-Isolated Genotypes of <i>Mycobacterium bovis</i> Vary in Virulence and Influence Case Pathology but Do Not Affect Outbreak Size. <i>PLoS ONE</i> , 2013, 8, e74503.	1.1	31
15	Genes controlling vaccine responses and disease resistance to respiratory viral pathogens in cattle. <i>Veterinary Immunology and Immunopathology</i> , 2012, 148, 90-99.	0.5	31
16	Living with the enemy or uninvited guests: Functional genomics approaches to investigating host resistance or tolerance traits to a protozoan parasite, <i>Theileria annulata</i> , in cattle. <i>Veterinary Immunology and Immunopathology</i> , 2012, 148, 178-189.	0.5	36
17	Adaptive evolution of Toll-like receptor 5 in domesticated mammals. <i>BMC Evolutionary Biology</i> , 2012, 12, 122.	3.2	38
18	The molecular pathways underlying host resistance and tolerance to pathogens. <i>Frontiers in Genetics</i> , 2012, 3, 263.	1.1	35

#	ARTICLE	IF	CITATIONS
19	Cytokine responses of Holstein and Sahiwal zebu derived monocytes after mycobacterial infection. <i>Tropical Animal Health and Production</i> , 2012, 44, 651-655.	0.5	10
20	Quantitative Trait Loci Associated with the Immune Response to a Bovine Respiratory Syncytial Virus Vaccine. <i>PLoS ONE</i> , 2012, 7, e33526.	1.1	19
21	Evidence for genetic variance in resistance to tuberculosis in Great Britain and Irish Holstein-Friesian populations. <i>BMC Proceedings</i> , 2011, 5, S15.	1.8	8
22	Strengthening insights into host responses to mastitis infection in ruminants by combining heterogeneous microarray data sources. <i>BMC Genomics</i> , 2011, 12, 225.	1.2	58
23	Quantitative trait loci for variation in immune response to a Foot-and-Mouth Disease virus peptide. <i>BMC Genetics</i> , 2010, 11, 107.	2.7	33
24	Bovine tuberculosis: the genetic basis of host susceptibility. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2010, 277, 2737-2745.	1.2	83
25	TGF- β 2 Induction Regulates Invasiveness of Theileria-Transformed Leukocytes and Disease Susceptibility. <i>PLoS Pathogens</i> , 2010, 6, e1001197.	2.1	62
26	Traits associated with innate and adaptive immunity in pigs: heritability and associations with performance under different health status conditions. <i>Genetics Selection Evolution</i> , 2009, 41, 54.	1.2	88
27	Comparative genomics of Toll-like receptor signalling in five species. <i>BMC Genomics</i> , 2009, 10, 216.	1.2	36
28	The protozoan parasite <i>Theileria annulata</i> alters the differentiation state of the infected macrophage and suppresses musculoaponeurotic fibrosarcoma oncogene (MAF) transcription factors. <i>International Journal for Parasitology</i> , 2009, 39, 1099-1108.	1.3	18
29	The Genome Sequence of Taurine Cattle: A Window to Ruminant Biology and Evolution. <i>Science</i> , 2009, 324, 522-528.	6.0	1,038
30	BoLA-DR peptide binding pockets are fundamental for foot-and-mouth disease virus vaccine design in cattle. <i>Vaccine</i> , 2009, 28, 28-37.	1.7	56
31	Molecular cloning and characterization of Toll-like receptors 1-10 in sheep. <i>Veterinary Immunology and Immunopathology</i> , 2009, 127, 94-105.	0.5	52
32	Variation matters: TLR structure and species-specific pathogen recognition. <i>Trends in Immunology</i> , 2009, 30, 124-130.	2.9	229
33	A rapid and robust sequence-based genotyping method for <i>BoLA-DRB3</i> alleles in large numbers of heterozygous cattle. <i>Animal Genetics</i> , 2008, 39, 561-563.	0.6	100
34	Molecular evolution of bovine Toll-like receptor 2 suggests substitutions of functional relevance. <i>BMC Evolutionary Biology</i> , 2008, 8, 288.	3.2	70
35	Differences in the transcriptional responses induced by <i>Theileria annulata</i> infection in bovine monocytes derived from resistant and susceptible cattle breeds. <i>International Journal for Parasitology</i> , 2008, 38, 313-325.	1.3	54
36	Review: Innate immunity to tropical theileriosis. <i>Innate Immunity</i> , 2008, 14, 5-12.	1.1	38

#	ARTICLE	IF	CITATIONS
37	Pig peripheral blood mononuclear leucocyte subsets are heritable and genetically correlated with performance. <i>Animal</i> , 2008, 2, 1575-1584.	1.3	42
38	The association between plasma levels of acute phase proteins, haptoglobin, alpha-1 acid glycoprotein (AGP), Pig-MAP, transthyretin and serum amyloid A (SAA) in Large White and Meishan pigs. <i>Veterinary Immunology and Immunopathology</i> , 2007, 119, 303-309.	0.5	24
39	Resistance and susceptibility to a protozoan parasite of cattle—Gene expression differences in macrophages from different breeds of cattle. <i>Veterinary Immunology and Immunopathology</i> , 2007, 120, 20-30.	0.5	51
40	Using genomic approaches to unravel livestock (host)–tick–pathogen interactions. <i>Trends in Parasitology</i> , 2007, 23, 439-444.	1.5	15
41	Analysis of the real EADGENE data set: Multivariate approaches and post analysis (Open Access) <i>Tj ETQq1 1 0.784314 rgBT /Overlo</i>	1.2	1
42	Quantitative evaluation of genetic and environmental parameters determining antibody response induced by vaccination against bovine respiratory syncytial virus. <i>Vaccine</i> , 2006, 24, 4007-4016.	1.7	30
43	Molecular immunophenotyping of lungs and spleens in naive and vaccinated chickens early after pulmonary avian influenza A (H9N2) virus infection. <i>Vaccine</i> , 2006, 24, 6096-6109.	1.7	34
44	Radiation hybrid mapping of all 10 characterized bovine Toll-like receptors. <i>Animal Genetics</i> , 2006, 37, 47-50.	0.6	83
45	Construction of a normalized <i>Bos taurus</i> and <i>Bos indicus</i> macrophage-specific cDNA library. <i>Animal Genetics</i> , 2006, 37, 75-77.	0.6	7
46	Development and validation of a bovine macrophage specific cDNA microarray. <i>BMC Genomics</i> , 2006, 7, 224.	1.2	36
47	Development of a chicken 5 K microarray targeted towards immune function. <i>BMC Genomics</i> , 2006, 7, 49.	1.2	32
48	Selection for lean growth and food intake leads to correlated changes in innate immune traits in Large White pigs. <i>Animal Science</i> , 2006, 82, 867-876.	1.3	12
49	<i>Bos taurus</i> and <i>Bos indicus</i> (Sahiwal) calves respond differently to infection with <i>Theileria annulata</i> and produce markedly different levels of acute phase proteins. <i>International Journal for Parasitology</i> , 2005, 35, 337-347.	1.3	118
50	Associations of weight gain and food intake with leukocyte sub-sets in Large White pigs. <i>Livestock Science</i> , 2005, 96, 249-260.	1.2	15
51	Innate immune traits differ between Meishan and Large White pigs. <i>Veterinary Immunology and Immunopathology</i> , 2005, 104, 131-144.	0.5	67
52	The expanding role of microarrays in the investigation of macrophage responses to pathogens. <i>Veterinary Immunology and Immunopathology</i> , 2005, 105, 259-275.	0.5	29
53	In-silico identification of chicken immune-related genes. <i>Immunogenetics</i> , 2004, 56, 122-133.	1.2	62
54	Genetic variation and responses to vaccines. <i>Animal Health Research Reviews</i> , 2004, 5, 197-208.	1.4	53

#	ARTICLE	IF	CITATIONS
55	Quantitative analysis of pro-inflammatory cytokine mRNA expression in <i>Theileria annulata</i> -infected cell lines derived from resistant and susceptible cattle. <i>Veterinary Immunology and Immunopathology</i> , 2004, 99, 87-98.	0.5	49
56	The protozoan parasite, <i>Theileria annulata</i> , induces a distinct acute phase protein response in cattle that is associated with pathology. <i>International Journal for Parasitology</i> , 2003, 33, 1409-1418.	1.3	51
57	Characterization of efferent lymph cells and their function following immunization of cattle with an allogenic <i>Theileria annulata</i> infected cell line. <i>Veterinary Immunology and Immunopathology</i> , 2003, 93, 39-49.	0.5	7
58	Functional expression of a bovine major histocompatibility complex class I gene in transgenic mice. <i>Veterinary Immunology and Immunopathology</i> , 2002, 87, 417-421.	0.5	2
59	The balance between protective immunity and pathogenesis in tropical theileriosis: what we need to know to design effective vaccines for the future. <i>Research in Veterinary Science</i> , 2001, 70, 71-75.	0.9	26
60	Proinflammatory cytokine expression by <i>Theileria annulata</i> infected cell lines correlates with the pathology they cause in vivo. <i>Vaccine</i> , 2001, 19, 2932-2944.	1.7	41
61	Sequence and transfection of BoLA β DRB3 cDNAs. <i>Animal Genetics</i> , 2000, 31, 219-222.	0.6	11
62	Reciprocal cross-protection induced by sporozoite antigens SPAG-1 from <i>Theileria annulata</i> and p67 from <i>Theileria parva</i> . <i>Parasite Immunology</i> , 2000, 22, 223-230.	0.7	22
63	Association of bovine DRB3 alleles with immune response to FMDV peptides and protection against viral challenge. <i>Vaccine</i> , 2000, 19, 1167-1171.	1.7	59
64	Evaluation of recombinant sporozoite antigen SPAG-1 as a vaccine candidate against <i>Theileria annulata</i> by the use of different delivery systems. <i>Tropical Medicine and International Health</i> , 1999, 4, A71-A77.	1.0	20
65	Comparative organization and function of the major histocompatibility complex of domesticated cattle. <i>Immunological Reviews</i> , 1999, 167, 145-158.	2.8	125
66	Innate and Adaptive Immune Responses Co-operate to Protect Cattle against <i>Theileria annulata</i> . <i>Parasitology Today</i> , 1999, 15, 268-274.	3.1	60
67	A Stage-specific, Parasite-induced, α Window of In Vivo Interferon- γ Production Is Associated with Pathogenesis in <i>Theileria annulata</i> Infection. <i>Annals of the New York Academy of Sciences</i> , 1998, 849, 152-154.	1.8	6
68	Different Vaccine Strategies Used to Protect against <i>Theileria annulata</i> . <i>Annals of the New York Academy of Sciences</i> , 1998, 849, 234-246.	1.8	43
69	Cytokine Production/T-Cell-stimulatory Ability of <i>Theileria annulata</i> -infected Cells and Post-Vaccinal Reactions. <i>Annals of the New York Academy of Sciences</i> , 1998, 849, 412-415.	1.8	4
70	Evidence for strain specificity in cytotoxic T-lymphocyte-mediated, major histocompatibility complex class I-dependent killing of <i>Theileria annulata</i> -infected cells. <i>Parasitology Research</i> , 1998, 84, 593-595.	0.6	19
71	Phenotypic and genotypic alterations associated with the attenuation of a <i>Theileria annulata</i> vaccine cell line from Turkey. <i>Vaccine</i> , 1998, 16, 569-575.	1.7	23
72	Parasite-Mediated Steps in Immune Response Failure During Primary <i>Theileria Annulata</i> Infection. <i>Tropical Animal Health and Production</i> , 1997, 29, 133S-135S.	0.5	4

#	ARTICLE	IF	CITATIONS
73	Characterization of cattle cDNA sequences from two DQA loci. <i>Immunogenetics</i> , 1997, 45, 455-458.	1.2	19
74	A non-protective T helper 1 response against the intra-macrophage protozoan <i>Theileria annulata</i> . <i>Clinical and Experimental Immunology</i> , 1997, 108, 463-470.	1.1	45
75	Functional expression of a cattle MHC class II DR-like antigen on mouse L cells. <i>Immunogenetics</i> , 1996, 43, 296-303.	1.2	4
76	<i>Theileria annulata</i> sporozoite antigen fused to hepatitis B core antigen used in a vaccination trial. <i>Vaccine</i> , 1995, 13, 1152-1160.	1.7	50
77	Bovine T cells preferentially recognize non-viral spacer epitopes in a putative FMDV vaccinal peptide. <i>Vaccine</i> , 1995, 13, 225-229.	1.7	22
78	<i>Theileria annulata</i> sporozoite targets. <i>Parasite Immunology</i> , 1994, 16, 501-505.	0.7	27
79	Parasite-accessory cell interactions in Theileriosis. Antigen presentation by <i>Theileria annulata</i> -infected macrophages and production of continuously growing antigen-presenting cell lines. <i>European Journal of Immunology</i> , 1990, 20, 2491-2497.	1.6	38
80	Generation and characterisation of bovine antigen-specific T cell lines. <i>Journal of Immunological Methods</i> , 1990, 128, 267-275.	0.6	15
81	Infection of bovine monocyte/macrophage populations with <i>Theileria annulata</i> and <i>Theileria parva</i> . <i>Veterinary Immunology and Immunopathology</i> , 1989, 22, 355-368.	0.5	59
82	Bovine mononuclear cell lines transformed by <i>Theileria parva</i> or <i>Theileria annulata</i> express different subpopulation markers. <i>Parasite Immunology</i> , 1988, 10, 619-629.	0.7	33
83	Qualitative variation in the immune response to ovarian follicular fluid proteins in cattle. <i>Journal of Reproductive Immunology</i> , 1988, 14, 151-163.	0.8	5
84	Impairment of monocyte ?lectin-like? receptor activity in Type 1 (insulin-dependent) diabetic patients. <i>Diabetologia</i> , 1987, 30, 228-231.	2.9	24
85	Modulation of mouse macrophage receptors by various inflammatory agents. <i>Agents and Actions</i> , 1984, 15, 12-13.	0.7	2
86	Macrophage binding of <i>Staphylococcus albus</i> is blocked by anti I-region alloantibody. <i>Nature</i> , 1982, 298, 852-854.	13.7	15