

Auriol C Purdie

List of Publications by Year in descending order

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Version: 2024-02-01

40
papers

1,280
citations

430874

18
h-index

361022

35
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43
all docs

43
docs citations

43
times ranked

1529
citing authors

#	ARTICLE	IF	CITATIONS
1	Leptospirosis is an emerging infectious disease of pig-hunting dogs and humans in North Queensland. PLoS Neglected Tropical Diseases, 2022, 16, e0010100.	3.0	12
2	Mycobacterium avium subsp. paratuberculosis exploits miRNA expression to modulate lipid metabolism and macrophage polarisation pathways during infection. Scientific Reports, 2022, 12, .	3.3	2
3	Mycobacterial infection-induced miR-206 inhibits protective neutrophil recruitment via the CXCL12/CXCR4 signalling axis. PLoS Pathogens, 2021, 17, e1009186.	4.7	18
4	Comparison of methods for miRNA isolation and quantification from ovine plasma. Scientific Reports, 2020, 10, 825.	3.3	52
5	The immunogenicity and tissue reactivity of Mycobacterium avium subsp paratuberculosis inactivated whole cell vaccine is dependent on the adjuvant used. Heliyon, 2019, 5, e01911.	3.2	11
6	The humoral immune response is essential for successful vaccine protection against paratuberculosis in sheep. BMC Veterinary Research, 2019, 15, 223.	1.9	18
7	Biomarkers for Detecting Resilience against Mycobacterial Disease in Animals. Infection and Immunity, 2019, 88, .	2.2	20
8	Gene expression profiles during subclinical Mycobacterium avium subspecies paratuberculosis infection in sheep can predict disease outcome. Scientific Reports, 2019, 9, 8245.	3.3	21
9	Mycobacterium avium subspecies paratuberculosis is able to manipulate host lipid metabolism and accumulate cholesterol within macrophages. Microbial Pathogenesis, 2019, 130, 44-53.	2.9	39
10	Immunopathological changes and apparent recovery from infection revealed in cattle in an experimental model of Johneâ€™s disease using a lyophilised culture of Mycobacterium avium subspecies paratuberculosis. Veterinary Microbiology, 2018, 219, 53-62.	1.9	22
11	Defining resilience to mycobacterial disease: Characteristics of survivors of ovine paratuberculosis. Veterinary Immunology and Immunopathology, 2018, 195, 56-64.	1.2	19
12	Analysis of mycobacterial infection-induced changes to host lipid metabolism in a zebrafish infection model reveals a conserved role for LDLR in infection susceptibility. Fish and Shellfish Immunology, 2018, 83, 238-242.	3.6	8
13	Integrated vaccine screening system: using cellular functional capacity in vitro to assess genuine vaccine protectiveness in ruminants. Pathogens and Disease, 2018, 76, .	2.0	5
14	Mycobacterium marinum infection drives foam cell differentiation in zebrafish infection models. Developmental and Comparative Immunology, 2018, 88, 169-172.	2.3	28
15	Sheep and cattle exposed to Mycobacterium avium subspecies paratuberculosis exhibit altered total serum cholesterol profiles during the early stages of infection. Veterinary Immunology and Immunopathology, 2018, 202, 164-171.	1.2	4
16	IFN-Î³ fails to overcome inhibition of selected macrophage activation events in response to pathogenic mycobacteria. PLoS ONE, 2017, 12, e0176400.	2.5	9
17	Evaluation of the limitations and methods to improve rapid phage-based detection of viable Mycobacterium avium subsp. paratuberculosis in the blood of experimentally infected cattle. BMC Veterinary Research, 2016, 12, 115.	1.9	14
18	A Rapid Method for Quantifying Viable Mycobacterium avium subsp. paratuberculosis in Cellular Infection Assays. Applied and Environmental Microbiology, 2016, 82, 5553-5562.	3.1	20

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19	Efficient, Validated Method for Detection of Mycobacterial Growth in Liquid Culture Media by Use of Bead Beating, Magnetic-Particle-Based Nucleic Acid Isolation, and Quantitative PCR. <i>Journal of Clinical Microbiology</i> , 2015, 53, 1121-1128.	3.9	22
20	Specific faecal antibody responses in sheep infected with <i>Mycobacterium avium</i> subspecies paratuberculosis. <i>Veterinary Immunology and Immunopathology</i> , 2015, 166, 125-131.	1.2	14
21	CD4+ T-cells, $\gamma\gamma$ T-cells and B-cells are associated with lack of vaccine protection in <i>Mycobacterium avium</i> subspecies paratuberculosis infection. <i>Vaccine</i> , 2015, 33, 149-155.	3.8	19
22	Development of 316v antibody enzyme-linked immunosorbent assay for detection of paratuberculosis in sheep. <i>OIE Revue Scientifique Et Technique</i> , 2015, 34, 869-879.	1.2	2
23	Cellular and humoral immune responses in sheep vaccinated with candidate antigens MAP2698c and MAP3567 from <i>Mycobacterium avium</i> subspecies paratuberculosis. <i>Frontiers in Cellular and Infection Microbiology</i> , 2014, 4, 93.	3.9	10
24	Antigenicity in sheep of synthetic peptides derived from stress-regulated <i>Mycobacterium avium</i> subsp. paratuberculosis proteins and comparison with recombinant protein and complex native antigens. <i>Veterinary Immunology and Immunopathology</i> , 2014, 158, 46-52.	1.2	2
25	Lymphoproliferative and Gamma Interferon Responses to Stress-Regulated <i>Mycobacterium avium</i> subsp. paratuberculosis Recombinant Proteins. <i>Vaccine Journal</i> , 2014, 21, 831-837.	3.1	8
26	High-Throughput Direct Fecal PCR Assay for Detection of <i>Mycobacterium avium</i> subsp. paratuberculosis in Sheep and Cattle. <i>Journal of Clinical Microbiology</i> , 2014, 52, 745-757.	3.9	76
27	Expression of genes associated with cholesterol and lipid metabolism identified as a novel pathway in the early pathogenesis of <i>Mycobacterium avium</i> subspecies paratuberculosis-infection in cattle. <i>Veterinary Immunology and Immunopathology</i> , 2014, 160, 147-157.	1.2	24
28	Immunoreactivity of protein tyrosine phosphatase A (PtpA) in sera from sheep infected with <i>Mycobacterium avium</i> subspecies paratuberculosis. <i>Veterinary Immunology and Immunopathology</i> , 2014, 160, 129-132.	1.2	4
29	Can early host responses to mycobacterial infection predict eventual disease outcomes?. <i>Preventive Veterinary Medicine</i> , 2013, 112, 203-212.	1.9	37
30	Antigenicity of Recombinant Maltose Binding Protein- <i>Mycobacterium avium</i> subsp. paratuberculosis Fusion Proteins with and without Factor Xa Cleaving. <i>Vaccine Journal</i> , 2013, 20, 1817-1826.	3.1	8
31	Development and Validation of a Liquid Medium (M7H9C) for Routine Culture of <i>Mycobacterium avium</i> subsp. paratuberculosis To Replace Modified Bactec 12B Medium. <i>Journal of Clinical Microbiology</i> , 2013, 51, 3993-4000.	3.9	52
32	<i>In Silico</i> Identification of Epitopes in <i>Mycobacterium avium</i> subsp. paratuberculosis Proteins That Were Upregulated under Stress Conditions. <i>Vaccine Journal</i> , 2012, 19, 855-864.	3.1	27
33	Comparative immunological and microbiological aspects of paratuberculosis as a model mycobacterial infection. <i>Veterinary Immunology and Immunopathology</i> , 2012, 148, 29-47.	1.2	310
34	Enhancement of the interferon gamma assay to detect paratuberculosis using interleukin-7 and interleukin-12 potentiation. <i>Veterinary Immunology and Immunopathology</i> , 2012, 149, 28-37.	1.2	11
35	In silico screened <i>Mycobacterium avium</i> subsp. paratuberculosis (MAP) recombinant proteins upregulated under stress conditions are immunogenic in sheep. <i>Veterinary Immunology and Immunopathology</i> , 2012, 149, 186-196.	1.2	12
36	Expression of genes associated with the antigen presentation and processing pathway are consistently regulated in early <i>Mycobacterium avium</i> subsp. paratuberculosis infection. <i>Comparative Immunology, Microbiology and Infectious Diseases</i> , 2012, 35, 151-162.	1.6	40

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37	Suppression of Avian Influenza Transmission in Genetically Modified Chickens. <i>Science</i> , 2011, 331, 223-226.	12.6	175
38	Indoleamine 2,3-Dioxygenase, Tryptophan Catabolism, and <i>Mycobacterium avium</i> subsp. <i>paratuberculosis</i> : a Model for Chronic Mycobacterial Infections. <i>Infection and Immunity</i> , 2011, 79, 3821-3832.	2.2	32
39	Candidate gene and genome-wide association studies of <i>Mycobacterium avium</i> subsp. <i>paratuberculosis</i> infection in cattle and sheep: A review. <i>Comparative Immunology, Microbiology and Infectious Diseases</i> , 2011, 34, 197-208.	1.6	45
40	Toll-like receptor (TLR)6 and TLR1 differentiation in gene expression studies of Johne's disease. <i>Veterinary Immunology and Immunopathology</i> , 2010, 137, 142-148.	1.2	23