

# Kumudika de Silva

## List of Publications by Year in descending order

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Version: 2024-02-01

52  
papers

1,855  
citations

331259

21  
h-index

264894

42  
g-index

56  
all docs

56  
docs citations

56  
times ranked

1527  
citing authors

#	ARTICLE	IF	CITATIONS
1	Mycobacterium avium subsp. paratuberculosis exploits miRNA expression to modulate lipid metabolism and macrophage polarisation pathways during infection. <i>Scientific Reports</i> , 2022, 12, .	1.6	2
2	Developing smarter vaccines for paratuberculosis: From early biomarkers to vaccine design. <i>Immunological Reviews</i> , 2021, 301, 145-156.	2.8	1
3	Mycobacterial infection-induced miR-206 inhibits protective neutrophil recruitment via the CXCL12/CXCR4 signalling axis. <i>PLoS Pathogens</i> , 2021, 17, e1009186.	2.1	18
4	Comparison of methods for miRNA isolation and quantification from ovine plasma. <i>Scientific Reports</i> , 2020, 10, 825.	1.6	52
5	Immunology of paratuberculosis infection and disease.. , 2020, , 248-265.		3
6	The immunogenicity and tissue reactivity of Mycobacterium avium subsp paratuberculosis inactivated whole cell vaccine is dependent on the adjuvant used. <i>Heliyon</i> , 2019, 5, e01911.	1.4	11
7	The humoral immune response is essential for successful vaccine protection against paratuberculosis in sheep. <i>BMC Veterinary Research</i> , 2019, 15, 223.	0.7	18
8	Biomarkers for Detecting Resilience against Mycobacterial Disease in Animals. <i>Infection and Immunity</i> , 2019, 88, .	1.0	20
9	Gene expression profiles during subclinical Mycobacterium avium subspecies paratuberculosis infection in sheep can predict disease outcome. <i>Scientific Reports</i> , 2019, 9, 8245.	1.6	21
10	Mycobacterium avium subspecies paratuberculosis is able to manipulate host lipid metabolism and accumulate cholesterol within macrophages. <i>Microbial Pathogenesis</i> , 2019, 130, 44-53.	1.3	39
11	Immunopathological changes and apparent recovery from infection revealed in cattle in an experimental model of Johne's disease using a lyophilised culture of Mycobacterium avium subspecies paratuberculosis. <i>Veterinary Microbiology</i> , 2018, 219, 53-62.	0.8	22
12	Defining resilience to mycobacterial disease: Characteristics of survivors of ovine paratuberculosis. <i>Veterinary Immunology and Immunopathology</i> , 2018, 195, 56-64.	0.5	19
13	Analysis of mycobacterial infection-induced changes to host lipid metabolism in a zebrafish infection model reveals a conserved role for LDLR in infection susceptibility. <i>Fish and Shellfish Immunology</i> , 2018, 83, 238-242.	1.6	8
14	Integrated vaccine screening system: using cellular functional capacity in vitro to assess genuine vaccine protectiveness in ruminants. <i>Pathogens and Disease</i> , 2018, 76, .	0.8	5
15	Sheep and cattle exposed to Mycobacterium avium subspecies paratuberculosis exhibit altered total serum cholesterol profiles during the early stages of infection. <i>Veterinary Immunology and Immunopathology</i> , 2018, 202, 164-171.	0.5	4
16	An objective method for assessment of foot conformation in sheep. <i>Small Ruminant Research</i> , 2018, 167, 22-28.	0.6	5
17	Immunological, clinical, haematological and oxidative responses to long distance transportation in horses. <i>Research in Veterinary Science</i> , 2017, 115, 78-87.	0.9	54
18	Applying the One Health Concept to Mycobacterial Research – Overcoming Parochialism. <i>Zoonoses and Public Health</i> , 2017, 64, 401-422.	0.9	16

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19	Variation in susceptibility of different breeds of sheep to <i>Mycobacterium avium</i> subspecies paratuberculosis following experimental inoculation. <i>Veterinary Research</i> , 2017, 48, 36.	1.1	18
20	Case definition terminology for paratuberculosis (Johne's disease). <i>BMC Veterinary Research</i> , 2017, 13, 328.	0.7	45
21	IFN- $\gamma$ fails to overcome inhibition of selected macrophage activation events in response to pathogenic mycobacteria. <i>PLoS ONE</i> , 2017, 12, e0176400.	1.1	9
22	Evaluation of the limitations and methods to improve rapid phage-based detection of viable <i>Mycobacterium avium</i> subsp. paratuberculosis in the blood of experimentally infected cattle. <i>BMC Veterinary Research</i> , 2016, 12, 115.	0.7	14
23	A Rapid Method for Quantifying Viable <i>Mycobacterium avium</i> subsp. paratuberculosis in Cellular Infection Assays. <i>Applied and Environmental Microbiology</i> , 2016, 82, 5553-5562.	1.4	20
24	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.	1.8	22
25	Specific faecal antibody responses in sheep infected with <i>Mycobacterium avium</i> subspecies paratuberculosis. <i>Veterinary Immunology and Immunopathology</i> , 2015, 166, 125-131.	0.5	14
26	Macrophage polarization in cattle experimentally exposed to <i>Mycobacterium avium</i> subsp. paratuberculosis. <i>Pathogens and Disease</i> , 2015, 73, ftv085.	0.8	41
27	CD4+ T-cells, $\gamma\delta$ 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.	1.7	19
28	Role of host- and pathogen-associated lipids in directing the immune response in mycobacterial infections, with emphasis on <i>Mycobacterium avium</i> subsp. paratuberculosis. <i>Critical Reviews in Microbiology</i> , 2014, 42, 1-13.	2.7	30
29	Lymphoproliferative and Gamma Interferon Responses to Stress-Regulated <i>Mycobacterium avium</i> subsp. paratuberculosis Recombinant Proteins. <i>Vaccine Journal</i> , 2014, 21, 831-837.	3.2	8
30	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.	1.8	76
31	Investigation of immunity in sheep following footrot infection and vaccination. <i>Vaccine</i> , 2014, 32, 6979-6985.	1.7	6
32	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.	0.5	24
33	Apoptosis of lymph node and peripheral blood cells in ovine Johne's disease. <i>Veterinary Immunology and Immunopathology</i> , 2013, 156, 82-90.	0.5	2
34	Can early host responses to mycobacterial infection predict eventual disease outcomes?. <i>Preventive Veterinary Medicine</i> , 2013, 112, 203-212.	0.7	37
35	Cellular and humoral immunogenicity of <i>Mycobacterium avium</i> subsp. paratuberculosis specific lipopentapeptide antigens. <i>Research in Veterinary Science</i> , 2013, 95, 123-129.	0.9	10
36	In vivo and in vitro expression pattern of Toll-like receptors in <i>Mycobacterium avium</i> subspecies paratuberculosis infection. <i>Veterinary Immunology and Immunopathology</i> , 2013, 156, 20-31.	0.5	17

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37	Development and Validation of a Liquid Medium (M7H9C) for Routine Culture of Mycobacterium avium subsp. paratuberculosis To Replace Modified Bactec 12B Medium. <i>Journal of Clinical Microbiology</i> , 2013, 51, 3993-4000.	1.8	52
38	Comparative immunological and microbiological aspects of paratuberculosis as a model mycobacterial infection. <i>Veterinary Immunology and Immunopathology</i> , 2012, 148, 29-47.	0.5	310
39	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.	0.5	11
40	Expression of genes associated with the antigen presentation and processing pathway are consistently regulated in early Mycobacterium avium subsp. paratuberculosis infection. <i>Comparative Immunology, Microbiology and Infectious Diseases</i> , 2012, 35, 151-162.	0.7	40
41	Does a Th1 over Th2 dominancy really exist in the early stages of Mycobacterium avium subspecies paratuberculosis infections?. <i>Immunobiology</i> , 2011, 216, 840-846.	0.8	376
42	Indoleamine 2,3-Dioxygenase, Tryptophan Catabolism, and Mycobacterium avium subsp. paratuberculosis: a Model for Chronic Mycobacterial Infections. <i>Infection and Immunity</i> , 2011, 79, 3821-3832.	1.0	32
43	The interleukin 10 response in ovine Johne's disease. <i>Veterinary Immunology and Immunopathology</i> , 2011, 139, 10-16.	0.5	29
44	Candidate gene and genome-wide association studies of Mycobacterium avium subsp. paratuberculosis infection in cattle and sheep: A review. <i>Comparative Immunology, Microbiology and Infectious Diseases</i> , 2011, 34, 197-208.	0.7	45
45	Experimental infection model for Johne's disease using a lyophilised, pure culture, seedstock of Mycobacterium avium subspecies paratuberculosis. <i>Veterinary Microbiology</i> , 2010, 141, 301-311.	0.8	57
46	The early lymphocyte proliferation response in sheep exposed to Mycobacterium avium subsp. paratuberculosis compared to infection status. <i>Immunobiology</i> , 2010, 215, 12-25.	0.8	32
47	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.	0.5	23
48	Enzyme-Linked Immunospot: An Alternative Method for the Detection of Interferon Gamma in Johne's Disease. <i>Journal of Veterinary Diagnostic Investigation</i> , 2009, 21, 187-196.	0.5	21
49	Identification of differentially expressed genes in ileum, intestinal lymph node and peripheral blood mononuclear cells of sheep infected with Mycobacterium avium subsp. paratuberculosis using differential display polymerase chain reaction. <i>Veterinary Immunology and Immunopathology</i> , 2009, 131, 177-189.	0.5	10
50	Toll-like receptor genes are differentially expressed at the sites of infection during the progression of Johne's disease in outbred sheep. <i>Veterinary Immunology and Immunopathology</i> , 2008, 124, 132-151.	0.5	47
51	Validation of endogenous reference genes for expression profiling of RAW264.7 cells infected with Mycobacterium avium subsp. paratuberculosis by quantitative PCR. <i>Veterinary Immunology and Immunopathology</i> , 2007, 115, 43-55.	0.5	13
52	Bone marrow stem cell and progenitor response to injury. <i>Wound Repair and Regeneration</i> , 2001, 9, 495-500.	1.5	21