

Kgomotso Sibeko-Matjila

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7726571/publications.pdf>

Version: 2024-02-01

21
papers

388
citations

1040056

9
h-index

752698

20
g-index

21
all docs

21
docs citations

21
times ranked

559
citing authors

#	ARTICLE	IF	CITATIONS
1	Development and evaluation of a real-time polymerase chain reaction test for the detection of <i>Theileria parva</i> infections in Cape buffalo (<i>Syncerus caffer</i>) and cattle. <i>Veterinary Parasitology</i> , 2008, 155, 37-48.	1.8	62
2	Molecular Detection of Zoonotic Rickettsiae and <i>Anaplasma</i> spp. in Domestic Dogs and Their Ectoparasites in Bushbuckridge, South Africa. <i>Vector-Borne and Zoonotic Diseases</i> , 2016, 16, 245-252.	1.5	52
3	Tick-borne haemoparasites in African buffalo (<i>Syncerus caffer</i>) from two wildlife areas in Northern Botswana. <i>Parasites and Vectors</i> , 2015, 8, 26.	2.5	50
4	Identification of <i>Theileria parva</i> and <i>Theileria</i> sp. (buffalo) 18S rRNA gene sequence variants in the African Buffalo (<i>Syncerus caffer</i>) in southern Africa. <i>Veterinary Parasitology</i> , 2011, 182, 150-162.	1.8	41
5	The epidemiology of tick-borne haemoparasites as determined by the reverse line blot hybridization assay in an intensively studied cohort of calves in western Kenya. <i>Veterinary Parasitology</i> , 2015, 210, 69-76.	1.8	41
6	Genetic analysis of the VP2-encoding gene of canine parvovirus strains from Africa. <i>Veterinary Microbiology</i> , 2013, 165, 460-465.	1.9	37
7	Four p67 alleles identified in South African <i>Theileria parva</i> field samples. <i>Veterinary Parasitology</i> , 2010, 167, 244-254.	1.8	26
8	Occurrence of tick-borne haemoparasites in cattle in the Mungwi District, Northern Province, Zambia. <i>Ticks and Tick-borne Diseases</i> , 2018, 9, 707-717.	2.7	13
9	Analyses of genes encoding <i>Theileria parva</i> p104 and polymorphic immunodominant molecule (PIM) reveal evidence of the presence of cattle-type alleles in the South African <i>T. parva</i> population. <i>Veterinary Parasitology</i> , 2011, 181, 120-130.	1.8	10
10	Selection and evaluation of housekeeping genes as endogenous controls for quantification of mRNA transcripts in <i>Theileria parva</i> using quantitative real-time polymerase chain reaction (qPCR). <i>PLoS ONE</i> , 2018, 13, e0196715.	2.5	10
11	Molecular detection and characterisation of protozoan and rickettsial pathogens in ticks from cattle in the pastoral area of Karamoja, Uganda. <i>Ticks and Tick-borne Diseases</i> , 2021, 12, 101709.	2.7	9
12	Influence of cycle stage, age and endometrial biopsy score on oxytocin receptor distribution and gene expression in the cervix and uterus of non-pregnant mares. <i>Theriogenology</i> , 2018, 120, 1-9.	2.1	8
13	Analysis of p67 allelic sequences reveals a subtype of allele type 1 unique to buffalo-derived <i>Theileria parva</i> parasites from southern Africa. <i>PLoS ONE</i> , 2020, 15, e0231434.	2.5	7
14	Genetic characterization of bovine viral diarrhoea (BVD) viruses: confirmation of the presence of BVD genotype 2 in Africa. <i>Archives of Virology</i> , 2013, 158, 155-163.	2.1	5
15	Molecular Analysis of South African Ovine Herpesvirus 2 Strains Based on Selected Glycoprotein and Tegument Genes. <i>PLoS ONE</i> , 2016, 11, e0147019.	2.5	4
16	Next generation sequencing and RNA-seq characterization of adipose tissue in the Nile crocodile (<i>Crocodylus niloticus</i>) in South Africa: Possible mechanism(s) of pathogenesis and pathophysiology of pansteatitis. <i>PLoS ONE</i> , 2019, 14, e0225073.	2.5	3
17	Microsatellite and minisatellite genotyping of <i>Theileria parva</i> population from southern Africa reveals possible discriminatory allele profiles with parasites from eastern Africa. <i>Ticks and Tick-borne Diseases</i> , 2020, 11, 101539.	2.7	3
18	South African Buffalo-Derived <i>Theileria parva</i> Is Distinct From Other Buffalo and Cattle-Derived <i>T. parva</i> . <i>Frontiers in Genetics</i> , 2021, 12, 666096.	2.3	3

#	ARTICLE	IF	CITATIONS
19	Unraveling the Complexity of the Rhomboid Serine Protease 4 Family of Babesia bovis Using Bioinformatics and Experimental Studies. Pathogens, 2022, 11, 344.	2.8	3
20	Whole genome sequencing of Theileria parva using target capture. Genomics, 2021, 113, 429-438.	2.9	1
21	Limited diversity in the CD8+ antigen-coding loci in Theileria parva parasites from cattle from southern and eastern Africa. Veterinary Parasitology, 2021, 291, 109371.	1.8	0