

Mikael Berg

List of Publications by Year in descending order

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Version: 2024-02-01

42
papers

810
citations

567144

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526166

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43
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43
docs citations

43
times ranked

1293
citing authors

#	ARTICLE	IF	CITATIONS
1	Crossing the Line: Seroprevalence and Risk Factors for Transboundary Animal Diseases Along the Tanzania-Zambia Border. <i>Frontiers in Veterinary Science</i> , 2022, 9, 809128.	0.9	2
2	Gizzard erosions in broiler chickens in Sweden caused by fowl adenovirus serotype 1 (FAdV-1): investigation of outbreaks, including whole-genome sequencing of an isolate. <i>Avian Pathology</i> , 2022, 51, 257-266.	0.8	4
3	Molecular Diversity of Hard Tick Species from Selected Areas of a Wildlife-Livestock Interface Ecosystem at Mikumi National Park, Morogoro Region, Tanzania. <i>Veterinary Sciences</i> , 2021, 8, 36.	0.6	4
4	Genetic Relationship Between Hard Ticks (Ixodidae) Infesting Cattle from Select Areas of a Wildlife-Livestock Interface Ecosystem at Mikumi National Park, Tanzania. <i>Vector-Borne and Zoonotic Diseases</i> , 2021, 21, 191-199.	0.6	1
5	Novel Viruses Found in Antricola Ticks Collected in Bat Caves in the Western Amazonia of Brazil. <i>Viruses</i> , 2020, 12, 48.	1.5	10
6	Field-Adapted Full Genome Sequencing of Peste-Des-Petits-Ruminants Virus Using Nanopore Sequencing. <i>Frontiers in Veterinary Science</i> , 2020, 7, 542724.	0.9	5
7	Genome constellations of rotavirus a isolated from avian species in Brazil, 2008-2015. <i>Brazilian Journal of Microbiology</i> , 2020, 51, 1363-1375.	0.8	4
8	Application of Viral Metagenomics for Study of Emerging and Reemerging Tick-Borne Viruses. <i>Vector-Borne and Zoonotic Diseases</i> , 2020, 20, 557-565.	0.6	8
9	Extinction and emergence of genomic haplotypes during the evolution of Avian coronavirus in chicken embryos. <i>Genetics and Molecular Biology</i> , 2020, 43, e20190064.	0.6	0
10	Complete Coding Sequence of a Pasivirus Found in Swedish Pigs. <i>Microbiology Resource Announcements</i> , 2020, 9, .	0.3	0
11	First-time detection of bovine viral diarrhoea virus, BVDV-1, in cattle in Botswana. <i>Onderstepoort Journal of Veterinary Research</i> , 2019, 86, e1-e7.	0.6	7
12	Updated unified phylogenetic classification system and revised nomenclature for Newcastle disease virus. <i>Infection, Genetics and Evolution</i> , 2019, 74, 103917.	1.0	227
13	Bornaviruses in naturally infected <i>Psittacus erithacus</i> in Portugal: insights of molecular epidemiology and ecology. <i>Infection Ecology and Epidemiology</i> , 2019, 9, 1685632.	0.5	0
14	Detection and phylogenetic analysis of parrot bornavirus 4 identified from a Swedish Blue-winged macaw (<i>Primolius maracana</i>) with unusual nonsuppurative myositis. <i>Infection Ecology and Epidemiology</i> , 2019, 9, 1547097.	0.5	3
15	Detection and genetic characterisation of porcine circovirus 3 from pigs in Sweden. <i>Virus Genes</i> , 2018, 54, 466-469.	0.7	74
16	Vector-borne viruses and their detection by viral metagenomics. <i>Infection Ecology and Epidemiology</i> , 2018, 8, 1553465.	0.5	3
17	Complete Genome Sequence of Avian Coronavirus Strain D274. <i>Microbiology Resource Announcements</i> , 2018, 7, .	0.3	1
18	Characterisation of the Virome of Tonsils from Conventional Pigs and from Specific Pathogen-Free Pigs. <i>Viruses</i> , 2018, 10, 382.	1.5	22

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19	Genetic characterization of a novel picorna-like virus in <i>Culex</i> spp. mosquitoes from Mozambique. <i>Virology Journal</i> , 2018, 15, 71.	1.4	21
20	Viral metagenomics reveals the presence of highly divergent quaranjavirus in <i>Rhipicephalus</i> ticks from Mozambique. <i>Infection Ecology and Epidemiology</i> , 2018, 8, 1478585.	0.5	20
21	Genome Sequence of a Bovine Rhinitis B Virus Identified in Cattle in Sweden. <i>Genome Announcements</i> , 2017, 5, .	0.8	5
22	Complete Genome of Avian coronavirus Vaccine Strains Ma5 and BR-I. <i>Genome Announcements</i> , 2017, 5, .	0.8	3
23	Seroprevalence and risk factors for peste des petits ruminants and selected differential diagnosis in sheep and goats in Tanzania. <i>Infection Ecology and Epidemiology</i> , 2017, 7, 1368336.	0.5	22
24	High seroprevalence of Rift Valley fever phlebovirus in domestic ruminants and African Buffaloes in Mozambique shows need for intensified surveillance. <i>Infection Ecology and Epidemiology</i> , 2017, 7, 1416248.	0.5	14
25	Single primer isothermal amplification (SPIA) combined with next generation sequencing provides complete bovine coronavirus genome coverage and higher sequence depth compared to sequence-independent single primer amplification (SISPA). <i>PLoS ONE</i> , 2017, 12, e0187780.	1.1	14
26	Viral Metagenomic Analysis Displays the Co-Infection Situation in Healthy and PMWS Affected Pigs. <i>PLoS ONE</i> , 2016, 11, e0166863.	1.1	34
27	History and current status of peste des petits ruminants virus in Tanzania. <i>Infection Ecology and Epidemiology</i> , 2016, 6, 32701.	0.5	24
28	Cloning, expression and characterization of potential immunogenic recombinant hemagglutinin-neuraminidase protein of Porcine rubulavirus. <i>Protein Expression and Purification</i> , 2016, 128, 1-7.	0.6	5
29	Seroprevalence of Rift Valley fever virus in sheep and goats in Zambia, Mozambique. <i>Infection Ecology and Epidemiology</i> , 2016, 6, 31343.	0.5	20
30	The Intestinal Eukaryotic Virome in Healthy and Diarrhoeic Neonatal Piglets. <i>PLoS ONE</i> , 2016, 11, e0151481.	1.1	28
31	Discovery of Novel Viruses in Mosquitoes from the Zambezi Valley of Mozambique. <i>PLoS ONE</i> , 2016, 11, e0162751.	1.1	42
32	Molecular and epidemiological studies of Porcine rubulavirus infection – an overview. <i>Infection Ecology and Epidemiology</i> , 2015, 5, 29602.	0.5	7
33	A longitudinal survey of African swine fever in Uganda reveals high apparent disease incidence rates in domestic pigs, but absence of detectable persistent virus infections in blood and serum. <i>BMC Veterinary Research</i> , 2015, 11, 106.	0.7	23
34	The origin of the PB1 segment of swine influenza A virus subtype H1N2 determines viral pathogenicity in mice. <i>Virus Research</i> , 2014, 188, 97-102.	1.1	6
35	Borna disease virus infection in cats. <i>Veterinary Journal</i> , 2014, 201, 142-149.	0.6	15
36	Isolation and characterization of low pathogenic H9N2 avian influenza A viruses from a healthy flock and its comparison to other H9N2 isolates. <i>Indian Journal of Virology: an Official Organ of Indian Virological Society</i> , 2013, 24, 342-348.	0.7	5

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37	The X proteins of bornaviruses interfere with type I interferon signalling. <i>Journal of General Virology</i> , 2013, 94, 263-269.	1.3	16
38	Markers of Borna disease virus infection in cats with staggering disease. <i>Journal of Feline Medicine and Surgery</i> , 2012, 14, 573-582.	0.6	13
39	Expression of interferon gamma in the brain of cats with natural Borna disease virus infection. <i>Veterinary Immunology and Immunopathology</i> , 2011, 141, 162-167.	0.5	9
40	Regulator of G protein signalling 16 is a target for a porcine circovirus type 2 protein. <i>Journal of General Virology</i> , 2009, 90, 2425-2436.	1.3	18
41	Development of a real-time RT-PCR assay for improved detection of Borna disease virus. <i>Journal of Virological Methods</i> , 2007, 143, 1-10.	1.0	21
42	Porcine circovirus type 2 replicase binds the capsid protein and an intermediate filament-like protein. <i>Journal of General Virology</i> , 2006, 87, 3215-3223.	1.3	50