Vincent Obanda

List of Publications by Year in descending order

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840776 713466 22 494 11 21 citations h-index g-index papers 23 23 23 1112 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Viral Population Diversity during Co-Infection of Foot-And-Mouth Disease Virus Serotypes SAT1 and SAT2 in African Buffalo in Kenya. Viruses, 2022, 14, 897.	3.3	4
2	Animal movement in a pastoralist population in the Maasai Mara Ecosystem in Kenya and implications for pathogen spread and control. Preventive Veterinary Medicine, 2021, 188, 105259.	1.9	6
3	Livestock Presence Influences the Seroprevalence of Crimean Congo Hemorrhagic Fever Virus on Sympatric Wildlife in Kenya. Vector-Borne and Zoonotic Diseases, 2021, 21, 809-816.	1.5	9
4	The role of African buffalo in the epidemiology of footâ€andâ€mouth disease in sympatric cattle and buffalo populations in Kenya. Transboundary and Emerging Diseases, 2020, 67, 2206.	3.0	14
5	The discovery and global distribution of novel mosquitoâ€associated viruses in the last decade (2007â€2017). Reviews in Medical Virology, 2019, 29, e2079.	8.3	48
6	Infection dynamics of gastrointestinal helminths in sympatric non-human primates, livestock and wild ruminants in Kenya. PLoS ONE, 2019, 14, e0217929.	2.5	14
7	Phylogeographical and crossâ€species transmission dynamics of SAT1 and SAT2 footâ€andâ€mouth disease virus in Eastern Africa. Molecular Ecology, 2019, 28, 2903-2916.	3.9	19
8	Molecular Detection and Genetic Characterization of Novel RNA Viruses in Wild and Synanthropic Rodents and Shrews in Kenya. Frontiers in Microbiology, 2019, 10, 2696.	3.5	16
9	Molecular identification of Ehrlichia, Anaplasma, Babesia and Theileria in African elephants and their ticks. PLoS ONE, 2019, 14, e0226083.	2.5	5
10	Adding injury to infection: The relationship between injury status and genetic diversity of Theileria infecting plains zebra, Equus quagga. Infection, Genetics and Evolution, 2018, 58, 269-278.	2.3	3
11	Metagenomic Virome Analysis of Culex Mosquitoes from Kenya and China. Viruses, 2018, 10, 30.	3.3	74
12	Molecular detection of viruses in Kenyan bats and discovery of novel astroviruses, caliciviruses and rotaviruses. Virologica Sinica, 2017, 32, 101-114.	3.0	54
13	Influence of Massive and Long Distance Migration on Parasite Epidemiology: Lessons from the Great Wildebeest Migration. EcoHealth, 2016, 13, 708-719.	2.0	12
14	The "strength of weak ties―and helminth parasitism in giraffe social networks. Behavioral Ecology, 2016, 27, 1190-1197.	2.2	33
15	Molecular Detection and Characterization of Theileria Infecting Wildebeest (Connochaetes taurinus) in the Maasai Mara National Reserve, Kenya. Pathogens, 2015, 4, 626-638.	2.8	7
16	Illegal tusk harvest and the decline of tusk size in the <scp>A</scp> frican elephant. Ecology and Evolution, 2015, 5, 5216-5229.	1.9	40
17	Epidemiology of Theileria bicornis among black and white rhinoceros metapopulation in Kenya. BMC Veterinary Research, 2015, 11, 4.	1.9	9
18	The Influence of Body Mass Index, Age and Sex on Inflammatory Disease Risk in Semi-Captive Chimpanzees. PLoS ONE, 2014, 9, e104602.	2.5	22

#	Article	IF	CITATION
19	Whole genome phylogenetic investigation of a West Nile virus strain isolated from a tick sampled from livestock in north eastern Kenya. Parasites and Vectors, 2014, 7, 542.	2.5	12
20	Mixed-host aggregations and helminth parasite sharing in an East African wildlife–livestock system. Veterinary Parasitology, 2014, 205, 224-232.	1.8	22
21	Isolation of Tick and Mosquito-Borne Arboviruses from Ticks Sampled from Livestock and Wild Animal Hosts in Ijara District, Kenya. Vector-Borne and Zoonotic Diseases, 2013, 13, 637-642.	1.5	53
22	Spatio-Temporal Distribution of Injured Elephants in Masai Mara and the Putative Negative and Positive Roles of the Local Community. PLoS ONE, 2013, 8, e71179.	2.5	18