

Benard Kulohoma

List of Publications by Year in descending order

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Version: 2024-02-01

14
papers

130
citations

1478458

6
h-index

1474186

9
g-index

15
all docs

15
docs citations

15
times ranked

225
citing authors

#	ARTICLE	IF	CITATIONS
1	Bone mineral density and risk of type 2 diabetes and coronary heart disease: A Mendelian randomization study. Wellcome Open Research, 2017, 2, 68.	1.8	26
2	Africa's response to the COVID-19 pandemic: A review of the nature of the virus, impacts and implications for preparedness. AAS Open Research, 0, 3, 19.	1.5	24
3	Kenya's response to the COVID-19 pandemic: a balance between minimising morbidity and adverse economic impact. AAS Open Research, 2021, 4, 3.	1.5	19
4	Prevalence of trypanosomes associated with drug resistance in Shimba Hills, Kwale County, Kenya. BMC Research Notes, 2020, 13, 234.	1.4	12
5	Antimicrobial Resistance Profiles and Genes of Staphylococci Isolated from Mastitic Cow's Milk in Kenya. Antibiotics, 2021, 10, 772.	3.7	9
6	Kenya's response to the COVID-19 pandemic: a balance between minimising morbidity and adverse economic impact. AAS Open Research, 2021, 4, 3.	1.5	8
7	Comparative genomics identifies male accessory gland proteins in five Glossina species. Wellcome Open Research, 2017, 2, 73.	1.8	8
8	Modelling appropriate use of trypanocides to restrict wide-spread multi-drug resistance during chemotherapy of animal African trypanosomiasis. Parasitology, 2019, 146, 774-780.	1.5	7
9	Comparative genomics identifies male accessory gland proteins in five Glossina species. Wellcome Open Research, 2017, 2, 73.	1.8	6
10	Community-led data collection using Open Data Kit for surveillance of animal African trypanosomiasis in Shimba hills, Kenya. BMC Research Notes, 2019, 12, 151.	1.4	4
11	Importance of human demographic history knowledge in genetic studies involving multi-ethnic cohorts. Wellcome Open Research, 2018, 3, 82.	1.8	3
12	H3Africa: crucial importance of knowledge on human demographic history in strategies for data exploitation – an analysis of the Luhya in Webuye, Kenya population from the 1000 Genomes Project. Wellcome Open Research, 2018, 3, 82.	1.8	3
13	Detecting bracoviral orthologs distribution in five tsetse fly species and the housefly genomes. BMC Research Notes, 2020, 13, 318.	1.4	1
14	NOMAD: metagenomic characterisation of the viral pathogen composition in outbreaks of non-malaria acute febrile illness cases. , 0, 5, 24.		0