

John B A Okello

List of Publications by Year in Descending Order

Source: <https://exaly.com/author-pdf/5214548/john-b-a-okello-publications-by-year.pdf>

Version: 2024-04-20

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

23
papers

553
citations

14
h-index

23
g-index

24
ext. papers

667
ext. citations

4.4
avg, IF

2.93
L-index

#	Paper	IF	Citations
23	GeneTerpret: a customizable multilayer approach to genomic variant prioritization and interpretation.. <i>BMC Medical Genomics</i> , 2022 , 15, 31	3.7	
22	Assessment of the Implementation of Pharmacogenomic Testing in a Pediatric Tertiary Care Setting. <i>JAMA Network Open</i> , 2021 , 4, e2110446	10.4	1
21	Risk Stratification of Prostate Cancer Through Quantitative Assessment of PTEN Loss (qPTEN). <i>Journal of the National Cancer Institute</i> , 2020 , 112, 1098-1104	9.7	12
20	The Cardiac Genome Clinic: implementing genome sequencing in pediatric heart disease. <i>Genetics in Medicine</i> , 2020 , 22, 1015-1024	8.1	15
19	Genes and Pathways Implicated in Tetralogy of Fallot Revealed by Ultra-Rare Variant Burden Analysis in 231 Genome Sequences. <i>Frontiers in Genetics</i> , 2020 , 11, 957	4.5	8
18	A three-gene DNA methylation biomarker accurately classifies early stage prostate cancer. <i>Prostate</i> , 2019 , 79, 1705-1714	4.2	19
17	Personalized risk stratification for patients with early prostate cancer (PRONTO): A Canadian team biomarker project.. <i>Journal of Clinical Oncology</i> , 2018 , 36, 109-109	2.2	
16	Reliability and performance of commercial RNA and DNA extraction kits for FFPE tissue cores. <i>PLoS ONE</i> , 2017 , 12, e0179732	3.7	32
15	Preparation of Formalin-fixed Paraffin-embedded Tissue Cores for both RNA and DNA Extraction. <i>Journal of Visualized Experiments</i> , 2016 ,	1.6	11
14	Genetic consequences of population expansions and contractions in the common hippopotamus (<i>Hippopotamus amphibius</i>) since the Late Pleistocene. <i>Molecular Ecology</i> , 2015 , 24, 2507-20	5.7	11
13	The episode of genetic drift defining the migration of humans out of Africa is derived from a large east African population size. <i>PLoS ONE</i> , 2014 , 9, e97674	3.7	17
12	Can small wildlife conservancies maintain genetically stable populations of large mammals? Evidence for increased genetic drift in geographically restricted populations of Cape buffalo in East Africa. <i>Molecular Ecology</i> , 2010 , 19, 1324-34	5.7	21
11	Quantitative assessment of the sensitivity of various commercial reverse transcriptases based on armored HIV RNA. <i>PLoS ONE</i> , 2010 , 5, e13931	3.7	23
10	Comparison of methods in the recovery of nucleic acids from archival formalin-fixed paraffin-embedded autopsy tissues. <i>Analytical Biochemistry</i> , 2010 , 400, 110-7	3.1	89
9	Where sociality and relatedness diverge: the genetic basis for hierarchical social organization in African elephants. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2009 , 276, 3513-21	4.4	52
8	Effective population size dynamics reveal impacts of historic climatic events and recent anthropogenic pressure in African elephants. <i>Molecular Ecology</i> , 2008 , 17, 3788-99	5.7	44
7	Mid-Holocene decline in African buffalos inferred from Bayesian coalescent-based analyses of microsatellites and mitochondrial DNA. <i>Molecular Ecology</i> , 2008 , 17, 4845-58	5.7	45

6	Population genetic structure of savannah elephants in Kenya: conservation and management implications. <i>Journal of Heredity</i> , 2008 , 99, 443-52	2.4	30
5	Age- and tactic-related paternity success in male African elephants. <i>Behavioral Ecology</i> , 2008 , 19, 9-15	2.3	50
4	Noninvasive genotyping and Mendelian analysis of microsatellites in African savannah elephants. <i>Journal of Heredity</i> , 2005 , 96, 679-87	2.4	24
3	Six new polymorphic microsatellite loci isolated and characterized from the African savannah elephant genome. <i>Molecular Ecology Notes</i> , 2005 , 5, 223-225		13
2	Mitochondrial DNA variation of the common hippopotamus: evidence for a recent population expansion. <i>Heredity</i> , 2005 , 95, 206-15	3.6	29
1	A recent bottleneck in the warthog and elephant populations of Queen Elizabeth National Park, revealed by a comparative study of four mammalian species in Uganda national parks. <i>Animal Conservation</i> , 2003 , 6, 237-245	3.2	7