

# Azza E Ahmed

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3045954/publications.pdf>

Version: 2024-02-01

13  
papers

175  
citations

1478505

6  
h-index

1474206

9  
g-index

15  
all docs

15  
docs citations

15  
times ranked

380  
citing authors

#	ARTICLE	IF	CITATIONS
1	Development of Bioinformatics Infrastructure for Genomics Research. Global Heart, 2017, 12, 91.	2.3	47
2	Ten simple rules for organizing a webinar series. PLoS Computational Biology, 2019, 15, e1006671.	3.2	43
3	Developing reproducible bioinformatics analysis workflows for heterogeneous computing environments to support African genomics. BMC Bioinformatics, 2018, 19, 457.	2.6	33
4	Organizing and running bioinformatics hackathons within Africa: The H3ABioNet cloud computing experience. AAS Open Research, 2018, 1, 9.	1.5	11
5	Organizing and running bioinformatics hackathons within Africa: The H3ABioNet cloud computing experience. AAS Open Research, 2018, 1, 9.	1.5	11
6	Delivering blended bioinformatics training in resource-limited settings: a case study on the University of Khartoum H3ABioNet node. Briefings in Bioinformatics, 2020, 21, 719-728.	6.5	9
7	Managing genomic variant calling workflows with Swift/T. PLoS ONE, 2019, 14, e0211608.	2.5	7
8	Design considerations for workflow management systems use in production genomics research and the clinic. Scientific Reports, 2021, 11, 21680.	3.3	7
9	A crowdsourced set of curated structural variants for the human genome. PLoS Computational Biology, 2020, 16, e1007933.	3.2	6
10	A crowdsourced set of curated structural variants for the human genome. , 2020, 16, e1007933.		0
11	A crowdsourced set of curated structural variants for the human genome. , 2020, 16, e1007933.		0
12	A crowdsourced set of curated structural variants for the human genome. , 2020, 16, e1007933.		0
13	A crowdsourced set of curated structural variants for the human genome. , 2020, 16, e1007933.		0