

Marion Adebisi

List of Publications by Year in descending order

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Version: 2024-02-01

25
papers

270
citations

1040056

9
h-index

940533

16
g-index

25
all docs

25
docs citations

25
times ranked

437
citing authors

#	ARTICLE	IF	CITATIONS
1	Strengthening Bioinformatics and Genomics Analyses Skills in Africa for Attainment of the Sustainable Development Goals: Report of the 2nd Conference of the Nigerian Bioinformatics and Genomics Network. American Journal of Tropical Medicine and Hygiene, 2022, , .	1.4	1
2	Windows Firewall Bypassing Techniques: An Overview of HTTP Tunneling and Nmap Evasion. Lecture Notes in Computer Science, 2021, , 546-556.	1.3	0
3	Eliminating Target Anopheles Proteins to Non-Target Organisms based on Posterior Probability Algorithm. Advances in Science, Technology and Engineering Systems, 2021, 6, 710-718.	0.5	0
4	The Nigerian Bioinformatics and Genomics Network (NBGN): a collaborative platform to advance bioinformatics and genomics in Nigeria. Global Health, Epidemiology and Genomics, 2020, 5, e3.	0.8	12
5	Severe Acne Skin Disease: A Fuzzy-Based Method for Diagnosis. Lecture Notes in Computer Science, 2020, , 320-334.	1.3	20
6	Diagmal: A Malaria Coactive Neuro-Fuzzy Expert System. Lecture Notes in Computer Science, 2020, , 428-441.	1.3	20
7	CAD-Based Machine Learning Project for Reducing Human-Factor-Related Errors in Medical Image Analysis. Advances in IT Personnel and Project Management, 2020, , 164-172.	0.3	5
8	Electronic Medical Information Encryption Using Modified Blowfish Algorithm. Lecture Notes in Computer Science, 2019, , 166-179.	1.3	12
9	A Deep Convolutional Encoder-Decoder Architecture for Retinal Blood Vessels Segmentation. Lecture Notes in Computer Science, 2019, , 180-189.	1.3	10
10	<p>An automated mammogram classification system using modified support vector machine</p>. Medical Devices: Evidence and Research, 2019, Volume 12, 275-284.	0.8	16
11	E-voting system: the pathway to free and fair election in Nigeria. Electronic Government, 2019, 15, 439.	0.2	4
12	Computational Investigation of Consistency and Performance of the Biochemical Network of the Malaria Parasite, Plasmodium falciparum. Lecture Notes in Computer Science, 2019, , 231-241.	1.3	4
13	Evaluation of the scholastic performance of students in 12 programs from a private university in the south-west geopolitical zone in Nigeria. F1000Research, 2019, 8, 154.	1.6	4
14	Evaluation of the scholastic performance of students in 12 programs from a private university in the south-west geopolitical zone in Nigeria. F1000Research, 2019, 8, 154.	1.6	6
15	Breathogenomics: A Computational Architecture for Screening, Early Diagnosis and Genotyping of Lung Cancer. Lecture Notes in Computer Science, 2017, , 41-49.	1.3	2
16	In Silico Prediction of 3D Structure of Anopheles Gambiae ABCC12 Protein. Lecture Notes in Computer Science, 2017, , 163-172.	1.3	0
17	Experimental Investigation of Frequency Chaos Game Representation for in Silico and Accurate Classification of Viral Pathogens from Genomic Sequences. Lecture Notes in Computer Science, 2017, , 155-164.	1.3	6
18	Assessing computational genomics skills: Our experience in the H3ABioNet African bioinformatics network. PLoS Computational Biology, 2017, 13, e1005419.	3.2	9

#	ARTICLE	IF	CITATIONS
19	Analyzing a single nucleotide polymorphism in schizophrenia: a meta-analysis approach. <i>Neuropsychiatric Disease and Treatment</i> , 2017, Volume 13, 2243-2250.	2.2	4
20	Development of Bioinformatics Infrastructure for Genomics Research. <i>Global Heart</i> , 2017, 12, 91.	2.3	47
21	Paving the Way Towards a Successful and Fulfilling Career in Computational Biology. <i>PLoS Computational Biology</i> , 2014, 10, e1003593.	3.2	4
22	Computational Biology and Bioinformatics in Nigeria. <i>PLoS Computational Biology</i> , 2014, 10, e1003516.	3.2	23
23	In Silico Models for Drug Resistance. <i>Methods in Molecular Biology</i> , 2013, 993, 39-65.	0.9	6
24	A Functional Workbench for <i>Anopheles gambiae</i> Micro Array Analysis. , 2013, , .		1
25	Ten Simple Rules for Organizing a Virtual Conference“Anywhere. <i>PLoS Computational Biology</i> , 2010, 6, e1000650.	3.2	54