

Oyediran Akinrinade

List of Publications by Year in descending order

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Version: 2024-02-01

21
papers

595
citations

686830

13
h-index

794141

19
g-index

25
all docs

25
docs citations

25
times ranked

1536
citing authors

#	ARTICLE	IF	CITATIONS
1	Genetics and genotype-phenotype correlations in Finnish patients with dilated cardiomyopathy. <i>European Heart Journal</i> , 2015, 36, 2327-2337.	1.0	130
2	Prevalence of Titin Truncating Variants in General Population. <i>PLoS ONE</i> , 2015, 10, e0145284.	1.1	85
3	Relevance of truncating titin mutations in dilated cardiomyopathy. <i>Clinical Genetics</i> , 2016, 90, 49-54.	1.0	43
4	GATA4 Is a Key Regulator of Steroidogenesis and Glycolysis in Mouse Leydig Cells. <i>Endocrinology</i> , 2015, 156, 1860-1872.	1.4	41
5	GATA4 Regulates Blood-Testis Barrier Function and Lactate Metabolism in Mouse Sertoli Cells. <i>Endocrinology</i> , 2016, 157, 2416-2431.	1.4	41
6	Loss of PPAR β in endothelial cells leads to impaired angiogenesis. <i>Journal of Cell Science</i> , 2016, 129, 693-705.	1.2	32
7	Relevance of Titin Missense and Non-Frameshifting Insertions/Deletions Variants in Dilated Cardiomyopathy. <i>Scientific Reports</i> , 2019, 9, 4093.	1.6	30
8	Copy Number Variation Analysis in Familial BRCA1/2-Negative Finnish Breast and Ovarian Cancer. <i>PLoS ONE</i> , 2013, 8, e71802.	1.1	26
9	Return of genetic and genomic research findings: experience of a pediatric biorepository. <i>BMC Medical Genomics</i> , 2019, 12, 173.	0.7	24
10	Quality of whole genome sequencing from blood versus saliva derived DNA in cardiac patients. <i>BMC Medical Genomics</i> , 2020, 13, 11.	0.7	23
11	Comparative genomics and genome biology of invasive <i>Campylobacter jejuni</i> . <i>Scientific Reports</i> , 2015, 5, 17300.	1.6	18
12	Machine Learning Identifies Clinical and Genetic Factors Associated With Anthracycline Cardiotoxicity in Pediatric Cancer Survivors. <i>JACC: CardioOncology</i> , 2020, 2, 690-706.	1.7	16
13	Germline copy number variation analysis in Finnish families with hereditary prostate cancer. <i>Prostate</i> , 2016, 76, 316-324.	1.2	14
14	Whole genome sequencing delineates regulatory, copy number, and cryptic splice variants in early onset cardiomyopathy. <i>Npj Genomic Medicine</i> , 2022, 7, 18.	1.7	14
15	Accessory genetic content in <i>Campylobacter jejuni</i> ST21CC isolates from feces and blood. <i>International Journal of Medical Microbiology</i> , 2017, 307, 233-240.	1.5	13
16	Transcription factor GATA4 associates with mesenchymal-like gene expression in human hepatoblastoma cells. <i>Tumor Biology</i> , 2018, 40, 101042831878549.	0.8	12
17	Everolimus Rescues the Phenotype of Elastin Insufficiency in Patient Induced Pluripotent Stem Cell-Derived Vascular Smooth Muscle Cells. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2020, 40, 1325-1339.	1.1	10
18	Targeted next generation sequencing reveals genetic defects underlying inherited retinal disease in Iranian families. <i>Molecular Vision</i> , 2019, 25, 106-117.	1.1	10

#	ARTICLE	IF	CITATIONS
19	Evolving Up-regulation of Biliary Fibrosis-Related Extracellular Matrix Molecules After Successful Portoenterostomy. <i>Hepatology Communications</i> , 2021, 5, 1036-1050.	2.0	7
20	Targeted oligonucleotide-selective sequencing of 101 genes from 150 patients with idiopathic dilated cardiomyopathy in Finland. <i>Heart Lung and Circulation</i> , 2014, 23, e2.	0.2	0
21	GENOMIC ARCHITECTURE OF TETRALOGY OF FALLOT AND GENOMIC PREDICTORS OF ADVERSE RIGHT VENTRICULAR REMODELLING. <i>Canadian Journal of Cardiology</i> , 2018, 34, S57.	0.8	0